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                                                                                               TELEFAX: 415-433-8716
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/164
FILING DATE: 08-DEC-1993
ATTORNEY/AGENT INFORMATION:
NAME: BOTTINEY, SCOLT R.
REGISTRATION NUMBER: 34,298
REFERENCE/DOCKET NUMBER: STRG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              109
                                                                                                                                                                                                                                                                                           SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                       HYPOTHETICAL:
                                  MOLECULE TYPE:
                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: 415-433-4150
                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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CITY: San Francisco
                                                           LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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                                                   TOPOLOGY:
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US-09-675-610B-24

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US-09-873-333D-1

US-09-539-333D-1

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HYPOTHETICAL: NO
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US-08-171-389-546/c
                                                                                                                                                                                                                            NAME: Fabian, Gary R.
REGISTRATION NUMBER: 3,875
REFERENCE/DOCKET NUMBER: 4600
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 324-0860
TELEPAX: (415) 324-0960
INFORMATION FOR SEQ ID NO: 546:
SEQUENCE CHARACTERISTICS:
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Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/081,070
FILING DATE: 22-JUN-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 07/996, OF PILLING DATE: 23-DEC-1992
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 07/723,618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA: PRIOR APPLICATION NUMBER: US 07/996,783
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TITLE OF INVENTION: Sequence-Directed DNA Binding
TITLE OF INVENTION: Molecules, Compositions and Methods
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MERAL TINEORUM... Edwards, Cynt... APPLICANT: ∠Edwards, Charles R. NoPLICANT: Cantor, Charles R. NoPLICANT: Andrews, Beth M.
                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 0: FILING DATE: 22-JUN-1993 ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: US 08/123,936
FILING DATE: 17-SEP-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Palan
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 505 Penobso
CITY: Redwood City
                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 0 FILING DATE: 27-JUN-1991
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                                                                                                                                                               TOPOLOGY:
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88.9%; Score 16; 100.0%; Pred. No.
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                                                                     Epstein Barr virus L2
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); Mismatches
DB 1; Length 50;
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RESULT 4
US-08-475-228A+546/c
; Sequence 546, Application US/08475228A
; Patent No. 5869241
                                                                                                                                                                                                                             ; INDIVIDUAL ISOLATE: Epstein Barr virus ; INDIVIDUAL ISOLATE: -90021) US-08-123-936-546
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US-08-123-936-546/c
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Best Local Similarity
GENERAL INFORMATION:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/996,783
FILING DATE: 23-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/723,618
APPLICATION NUMBER: US 07/723,618
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Fabian, Gary R.
REGISTRATION NUMBER: 33,875
REFERENCE/DOCKET NUMBER: 4600-0075.32/G19P2
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 324-0880
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Genelabs
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ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
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505 Penobscot Drive
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                                                                                                                                                                   Conservative
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Turin, Lisa M.
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Pred. No.
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RESULT 5

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US-08-482-080A-546/c

Sequence 546, Application US/08482080A Patent No. 6010849
GENERAL INFORMATION:

APPLICANT:

Edwards, Cynthia A. Cantor, Charles R. Andrews, Beth M. Turin, Lisa M.

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                                                                 Matches
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Best Local S
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                              ORIGINAL SOURCE: Epstein Barr virus L2
INDIVIDUAL ISOLATE: -90021)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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TITLE OF INVENTION: Sequence-Directed DNA Binding
TITLE OF INVENTION: Molecules, Compositions and Methods
                                                                                                                                                                                                                 MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 34,444
REFERENCE/DOCKET NUMBER: 4600-0175.21/G19P3D2
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 324-0880
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APPLICATION NUMBER: US/
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                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 22-JUN-1993
ATTORNEY/AGENT INFORMATION:
NAME: Stratford, Carol A.
                                                                                                                                                                                                                                                STRANDEDNESS:
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                                                                                                                                                                                                                                                              TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
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41 GGCTGGTGTCACCTGT 26
                         1 GGCTGGTGTCACCTGT 16
                                                               Similarity
16; Conserv
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                                                               Conservative
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Turin, Lisa M.
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                                                            100.0%; 1
                                                               88.9%; Score 16; DB
100.0%; Pred. No. 1.
Live 0; Mismatches
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                                                                                                                                       RESULT 6
                                                                          Patent No. 6384208
GENERAL INFORMATION:
                                                                                                         Sequence 546,
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/482,080A
FILING DATE: 07-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/171,389
FILING DATE: 20-DEC-1993
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 08/123,936
EILING DATE: 17-SEP-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/996,783
EILING DATE: 23-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/723,618
                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (650) 324-0960 INFORMATION FOR SEQ ID NO: 5 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                        ORIGINAL-SOURCE:
INDIVIDUAL-SOURCE: Epstein Barr virus L2
INDIVIDUAL ISOLATE: 90021)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (650) 324-0960
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRILING DATE: 27-JUN-
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Genelabs Technologies, Inc
                               APPLICANT:
                                            APPLICANT:
                                                            APPLICANT:
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TITLE OF INVENTION: Sequence-Directed DNA Binding
TITLE OF INVENTION: Molecules, Compositions and Methods
   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE:
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CITY: Redwood City
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                                                                                                                                                                                                                                                             Local
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16; Conservative
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                                                                                                         Application US/09354947
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Andrews, Beth M.
Turin, Lisa M.
Fry, Kirk E.
                                            Edwards, Cynthia A. Cantor, Charles R.
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                                                         Edwards,
                                                                                                                                                                                                                                                                                                                                                                                                     linear
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100.0%; Pr
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Pred. No.
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RESULT 7
PCT-US93-12388-546/c
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                                                                               Sequence 546, Application PC/TUS9312388 GENERAL INFORMATION:
                                                                                                                                                                                                                                                            Matches
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Best Local Similarity
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          TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE:
HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 50 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 46
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 324-0880
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Sequence-Directed DY
TITLE OF INVENTION: Molecules, Compositi
NUMBER OF SEQUENCES: 664
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genelabs Technologies, Inc.
STREET: 505 Penobscot Drive
                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
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PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 22-JUN-1993
ATTORNEY/AGENT INFORMATION:
NAME: Brady, John F.
REGISTRATION NUMBER: 39,118
BREGERIC/FORTET NIMBER: 36/1/8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                               INDIVIDUAL ISOLATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: FILING DATE: 23-DE0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Redwood City
                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 20-DEC-1993
                                                                                                                                                                                       41 GGCTGTGTCACCTGT 26
                                                                                                                                                                                                                                                          16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    nucleic acid
DEDNESS: double
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                                                                                                                                                                                                           GGCTGGTGTCACCTGT 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  50 base pairs
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                                                                                                                                                                                                                                                            Conservative
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23-DEC-1992
                                                                                                                                                                                                                                                                                                                                                                                                                DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-SEP-1993
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                           Sequence-Directed DNA Binding Molecules, Compositions and Methods
                                                                                                                                                                                                                                                                          88.9%; Score 16;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Release #1.0, Version #1.25
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                                                                                                                                                                                                                                                                                                                                               Epstein Barr virus L2 90021)
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ADDRESSEE: Genelabs Technologies, Inc.

505 Penobscot Drive

CLASSIFICATION:

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INDIVIDUAL ISOLATE:
CT-US93-12388-546
                                                                                                                                                                                                                                                                                                           Sequence 18, Application US/08917320 Patent No. 5824508
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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Best Local
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[NFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OPERATING SYSTEM: PC-DOS/MS-DOS
SOTTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/12388
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ORIGINAL SOURCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US 0
FILING DATE: 17-SEP-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
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        SOFTWARE: Patent In Re-
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
                                                                                           COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                            APPLICANT: Spaete, Richard and Jackman, Winthrop, TITLE OF INVENTION: No. 5824508 Splicing Variants NUMBER OF SEQUENCES: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLICE TYPE: UN
                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 324-0880
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
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TTORNEY/AGENT INFORMATION:
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OPERATING SYSTEM:
SOFTWARE: PatentI
                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                       STREET: 5 Paio -
CITY: Palo Alto
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REGISTRATION NUMBER: 33
REFERENCE/DOCKET NUMBER:
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CLASSIFICATION:
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CITY: Redwood City
                                                                                                                                            COUNTRY:
                                                                                                                                                                                              ADDRESSEE: Cooley Godward Castro Huddleson & Tatum
STREET: 5 Palo Alto Square
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                                                                                                                                                                                                                                                                                             INFORMATION:
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                                            Patent In Release #1.0, Version #1.25
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25-AUG-1997
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                                                                                                                                                                                                                                              No. 5824508 Splicing Variants of gp350/220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Epstein Barr virus L2 (start site 90021)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US 08/123,936
            US/08/917,320
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; LOCATION:
US-08-917-320-18
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Best Local Similarity
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                                                               INFORMATION FOR SEQ ID NO:
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LENGTH: 3833 base pair
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                                                                                                                              NAME: LUADN CSETE
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: AV
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                              SOPTWARE: Patent In Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/0461
                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE:
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                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/229,
FILING DATE: April 18, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Spaete, Richard and Jackman, Winthrop, T.
TITLE OF INVENTION: Non Splicing Variants of gp350/220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: USA
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STATE: California
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                                                                                            TELEFAX: 415-857-0663
                                                                                                              TELEPHONE:
                                                                                                                                                                                                                  CLASSIFICATION:
                                                                                                                                                                                                                                                                                    CLASSIFICATION:
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                            ENGTH:
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al Similarity 100.0%;
16; Conservative (
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        3833 base pairs
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; Pred. No.
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FEATURE

MOLECULE TYPE: CDNA

TOPOLOGY: unknown

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Matches
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Patent No. 6054130
GENERAL INFORMATION:
                                                  Matches
                                                                          Query Match
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                                                                                                                                                                                                                                           NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: FASTSEQ Version 2.0 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/783,774
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                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                               TELEPHONE: 212-790-9090
                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                 OLECULE TYPE:
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                                                                                                                                                                                                                                                        TELEPHONE: Z14
TELEPHONE: Z14
TENAX: 212-869-8864
                                                                                                                                                                                                                                                                                                       NAME: Coruzzi, Laura A.
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 76
                                                              Local
                                                                                                               NAME/KEY: Coding Sequence LOCATION: 1014...3734 OTHER INFORMATION:
                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: FILING DATE: 15-JA
                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
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3137
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                                               16;
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                       1 GGCTGGTGTCACCTGT 16
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                                                              Similarity
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                                                                                                                                                                                                                                                      66141 PENNIE
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                                                                                                                                                                                                                 5931 base pairs
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                                                  Conservative
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Jackman, Winthrop
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Diskette
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100.0%; Pr
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                                                              100.0%;
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                                                 Mismatches
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RESULT 11

; Sequence 3, Application US/09171710 Patent No. 6323330 ; GENERAL INFORMATION: APPLICANT: ISHIDUKA, Yasuyuki

US-09-171-710-3

RESULT 13

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                                                                                                                                                 ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-171-710-5
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US-09-171-710-5
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                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/171,710, CURRENT FILING DATE: 1998-10-23
EARLIER APPLICATION NUMBER: PCT/JP97/01391
EARLIER FILING DATE: 1997-10-30
EARLIER APPLICATION NUMBER: 9-41562
EARLIER FILING DATE: 1997-02-10
EARLIER APPLICATION NUMBER: 8-127954
EARLIER FILING DATE: 1996-04-23
                                                                                                                                                                                                                          SOFTWARE: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 1, Application US/09556706B Patent No. 6458364 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 5, Application Patent No. 6323330
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LENGTH: 5931
                                                                          Matches
                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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APPLICANT: Jackman, Winthrop
TITLE OF INVENTION: NON SPLICING VARIANTS OF GP350/220
FILE REFERENCE: 7682-050-99
FILE REFERENCE: 7682-050-99
CURRENT APPLICATION NUMBER: US/09/556,706B
CURRENT FILING DATE: 2000-04-24
PRIOR APPLICATION NUMBER: 08/783,774
PRIOR APPLICATION NUMBER: 08/783,774
PRIOR TILING DATE: 1997-01-15
PRIOR APPLICATION NUMBER: 08/229,291
RRIOR APPLICATION NUMBER: 08/229,291
RRIOR APPLICATION NUMBER: 08/229,291
RRIOR FILING DATE: 1994-04-18
                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: ISHIDUKA, Yasuyuki
APPLICANT: MOCHIZUKI, Reiko
TITLE OF INVENTION: NOVEL PROTEINS C16 AND C16N OR GENES ENCODING THE SAME
FILE REFERENCE: 0020-4474P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Virus FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
                                                                                                                                                                                                        LENGTH: 1808
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949 GGCTGGTGTCACCTG 963
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                       1 GGCTGGTGTCACCTG 15
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                                                                          Conservative
                                                                83.3%; ---
100.0%; Pr
0;
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100.0%; Pr/
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                                                                                          Score 15; DB 4;
Pred. No. 3.8;
                                                                          Mismatches
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0.99;
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US-07-745-206A-12
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Best Local
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TYPE: DNA
ORGANISM: Mus sp.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EARLIER APPLICATION NUMBER: 8-127954
EARLIER FILING DATE: 1996-04-23
NUMBER OF SEQ ID NOS: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/171,710 CURRENT FILING DATE: 1998-10-23 EARLIER APPLICATION NUMBER: PCT/JP97/01391 EARLIER FILING DATE: 1997-10-30 EARLIER APPLICATION NUMBER: 9-41562 EARLIER FILING DATE: 1997-02-10
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                                                                                                                                                                             NAME: Feder, Scott B
REFERENCE/DOCKET NUMBER: 51
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-372-7842
INFORMATION FOR SEQ ID NO: 12:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ILE REFERENCE: 0020-4474P
                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/745,206A
                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 5467 base pairs
TYPE: NUCLEIC ACID
                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                 NAME: Feder, Scott B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
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                                                                      OLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             UMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PPLICANT: Feldman, Daniel
ITLE OF INVENTION: Human Calcium Channel Compositions and
ITLE OF INVENTION: Methods
 NAME/KEY:
LOCATION:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                       STRANDEDNESS:
                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE:
                                                                                          TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OF INVENTION: NOVEL PROTEINS C16 AND C16N OR GENES ENCODING THE SAME
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15; Conserv
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135 S. LaSalle
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Williams, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               McCue, Ann
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                                                                                          unknown
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join(144..3164, 3168..3245, 3249..3386, 3390
..3392, 3396..3488, 3495..3539, 3543..3581, 3585
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                                                                                                           unknown
                                                                      DNA (genomic)
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100.0%; Pr
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k; Pred. No. 3.7
0; Mismatches
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3.7;
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US-07-745-206A-12
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                 Query Match
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ZIP: 92101-2926

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

MEDIUM TYPE: Floppy disk

PC-DOS/MS-DOS

MS-DOS

MS-DOS
                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 07/745,206
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compa
OPERATING SYSTEM: PC-D
SOFTWARE: PatentIn Rel
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 63
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                          FILING DATE: 15-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: Seidman, Stephanie L.
NAME: Seidman, Stephanie T.
REGISTRATION NUMBER: 33,779
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION:
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                                                                                                                                                                           MOLECULE TYPE:
Local
                                                                                                                             NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: San Diego
STATE: California
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                                                                                                                                                                                                        STRANDEDNESS:
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                                                                                                                LOCATION:
                                                                                                                                                                                                                                                                                                   TELEPHONE:
                                                                                                                                                                                            COPOLOGY:
                                                                                                                                                                                                                     YPE: nucleic acid
                                                                                                                                                                                                                                       ENGTH:
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 Similarity
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                                                                                                                                                                                                                                    5467 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Brenner, Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Williams, Mark
Feldman, Daniel
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..3746, 3750..4823, 4827..4841, 4845..5006, 
..5096, 5100..5306, 5310..5366, 5370..5465)
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                                                               .3392, 3396..3488, 3495..3539, 3543..3581, 3585
.3587, 3591..3626, 3600..3689, 3693..3737, 3744
..3746, 3750..4823, 4827..4841, 4845..5006, 5010
..5096, 5100..5306, 5310..5366, 5370..5465)
                                                                                                                           join(144..3164, 3168..3245, 3249..3386, 3390
                                                                                                                                                                                            unknown
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                                                                                                                                                                         DNA (genomic)
                                                                                                                                                                                                                                                                                                     (619) 238-0999
                                                                                                                                                                                                          unknown
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100.0%; F1
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 100.0%;
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 77.8%; Score 14; DB 2; Length 5467; 100.0%; Pred. No. 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                            US/08/311,363
                                                                                                                                                                                                                                                                                                                                    6362-51506
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14;
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; LOCATION: (148)...(177)
US-09-657-453A-24
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US-09-657-453A-24/c
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LENGTH: 261
TYPE: DNA
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Best Local Similarity
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CURRENT FILING DATE: 2000-09-07
NUMBER OF SEQ ID NOS: 105
                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/141,027A CURRENT FILING DATE: 1998-08-27 EARLIER APPLICATION NUMBER: 60/656,248 EARLIER FILING DATE: 1997-08-29
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TITLE OF INVENTION: ANT
FILE REFERENCE: RTS-013
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             equence 9, Application US/09141027A atent No. 6372454
                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 19
SOFTWARE: PatentIn Ver. :
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                                                                                                                     NAME/KEY: misc feature
LOCATION: (24) -
OTHER INFORMATION: n equals
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                                                           LOCATION: (29)
OTHER INFORMATION: n equals
                                                                                     NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                              ORGANISM: Homo sapiens FEATURE:
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OTHER INFORMATION: n equals a,
                NAME/KEY: misc_feature = LOCATION: (34)
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LOCATION: (3)
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RESULT 18 US-09-617-804-9/c

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Sequence 9, Application US/09617804 Patent No. 6537966 GENERAL INFORMATION:

APPLICANT: Duan, et al FILE REFERENCE:

PF388P1

Follistatin-3

CURRENT APPLICATION NUMBER: US/09/617,804
CURRENT FILING DATE: 2000-07-14
PRIOR APPLICATION NUMBER: 60/144,088
PRIOR FILING DATE: 199-07-16
PRIOR APPLICATION NUMBER: 09/141,027

FILING DATE:

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                                                                                 ; OTHER INFORMATION: n equals US-09-141-027-9
                                        Matches
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Best Local Similarity
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NAME/KEY: misc feature
COLITION: (297)
NAME/KEY: misc feature
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LOCATION: (308)
                                                                                                                                   OTHER INFORMATION: n equals
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LOCATION: (237)
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OTHER INFORMATION: n equals
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PRIOR FILING DATE: 1997-08-29
PRIOR APPLICATION NUMBER: PCT,
PRIOR FILING DATE: 1998-08-27
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                                                                       OTHER INFORMATION: n equals a,
                                                                                   NAME/KEY: misc feature LOCATION: (308)
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LOCATION: (297)
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; LOCATION: (1)...(361)
; OTHER INFORMATION: n = A,T,C or US-09-702-705-1159
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SOFTWARE: FastSEQ for
SEQ ID NO 1159
                                                                                                                     CURRENT APPLICATION NUMBER: US/09/736,457
CURRENT FILING DATE: 2000-12-13
NUMBER OF SEQ ID NOS: 1864
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 1159
LENGTH: 361
TYPE: DNA
TYPE: DNA
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Best Local Similarity
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Patent No.
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Query Match
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APPLICANT: Wang, Aijun
TITLE OF INVENTION: COMPOSITIONS AND METHODS
TITLE OF INVENTION: DIAGNOSIS OF LUNG CANCER
FILE REFERENCE: 210121.478C15
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CURRENT FILING DATE: 2000-10-30
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TITLE OF INVENTION: COMPOSITIONS AND METHODS
TITLE OF INVENTION: DIAGNOSIS OF LUNG CANCER
FILE REFERENCE: 210121.478C14
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                                              OTHER INFORMATION: n =
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Retter, Marc
Mannion, Jane
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Vedvick, Tom
Carter, Darrick
Retter, Marc
Mannion, Jane
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Lodes, Michael
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Lodes, Michael
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Length 361;
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; SEQ ID NO 10
; LENCTH: 794
; TYPE: DNA
; ORGANISM: Zea mays
US-09-173-300-10
              RESULT 23
US-09-221-456-1/c
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US-09-173-300-10
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Sequence 1, Application US/09221456
                                                                                                                                            Matches
                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Falco, Saverio Carl
APPLICANT: Hitz, William D.
APPLICANT: Kinney, Anthony J.
APPLICANT: Cahoon, Rebecca B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 10, Application US/09173300 Patent No. 6451581
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Patent No. 635919
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                                                                                                                                                                                                                                                                                                              FILE REFERENCE: BB-1126
CURRENT APPLICATION NUMBER: US/09/173,300
CURRENT FILING DATE: 1998-10-15
EARLIER APPLICATION NUMBER: 60/063,423
EARLIER FILING DATE: 1997 October 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/228,986
CURRENT FILLING DATE: 1999-01-12
NUMBER OF SEQ ID NOS: 130
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Compositions Isolated from Plant Cells
TITLE OF INVENTION: and Their Use in the Modification of Plant Cell Signalling
FILE REFERENCE: 11000/1020
                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Rafalski, J. Antoni
TITLE OF INVENTION: PLANT BRANCHED CHAIN AMINO ACID BIOSYNTHETIC ENZYMES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Strabala, Timothy APPLICANT: Nieuwenhuizen, Niels
                                                                                                                                                                                                                                                                                  SOFTWARE: Microsoft Word Version 7.0A
                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Pinus radiata
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; MOLECULE TYPE:
US-09-221-456-1
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Best Local (
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                 CURRENT APPLICATION NUMBER: US/09/558,740
CURRENT FILING DATE: 2000-04-26
PRIOR APPLICATION NUMBER: 08/956,975
PRIOR FILING DATE: 1997-10-23
                                                                                     APPLICANT: MUIR, ALISON
APPLICANT: CHAMBERS, JON
APPLICANT: SZEKERES, PHILIP
TITLE OF INVENTION: METHODS OF SCREENING FOR AGONISTS AND
TITLE OF INVENTION: ANTAGONISTS OF THE HNEAA81 RECEPTOR
FILE REFERENCE: GH-70318-2
                                                                                                                                                                                                  APPLICANT: HALSEY, WENDY
                                                                                                                                                                                                                   APPLICANT: SATHE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
RIOR APPLICATION NUMBER: 09/221,456
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/956,975
FILING DATE: 23-OCT-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    - ZIP: 19482
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: 1BM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
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CORRESPONDENCE ADDRESS:
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TITLE OF INVENTION: AND ANTAGONISTS OF THE HNEAA81 RECEPTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity 100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: 610-407-0700
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REFERENCE/DOCKET NUMBER: GH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Prestia, Paul F
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 1124 base pairs
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HALSEY, WENDY
MUIR, ALISON
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SZEKERES, PHILIP
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RESULT 26
US-09-328-352-2907/c
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                                                                                                                  ; ORGANISM: Acinetobacter baumannii US-09-328-352-2907
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APPLICANT: Gary L. Breton et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
TITLE OF INVENTION: BAUWANNII FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                               NUMBER OF SEQ ID NOS: 8252
SEQ ID NO 2907
LENGTH: 1716
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/328,352
CURRENT FILING DATE: 1999-06-04
NUMBER OF SEQ ID NOS: 8252
SEQ ID NO 3822
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Best Local Similarity
Matches 13; Conserv
                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: GATY L. Breton et al.
APPLICANT: GATY L. BRETON et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
TITLE OF INVENTION: BAUMANNII FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                           Matches
                                                                                                                                                                                                                                                                                                         Patent No. 656295
                                                                                                                                                                                                                                                                                                                                Sequence 2907,
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                                                                        Query Match
Best Local Similarity
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                                                                                                                                                                                                                         FILE REFERENCE: GTC99-03PA
CURRENT APPLICATION NUMBER: US/09/328,352
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                              CURRENT FILING DATE: 1999-06-04
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TYPE: DNA
ORGANISM: HOMO SAPIENS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA ORGANISM: Acinetobacter baumannii
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ENGTH: 1302
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1328
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                            6 GTGTCACCTGTTA 18
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GTGTCACCTGTTA 1316
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                                                          Conservative
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100.0%;
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Pred. No.
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Pred. No.
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53;
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52;
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RESULT 27

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                                                                                                                                                                Patent No.
                                                                                                                                                                          Sequence 2,
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                                                                                                                             GENERAL INFOR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM: Windows Version SOFTWARE: FastSEQ for Windows Version CURRENT APPLICATION DATA:
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MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                     IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Cerrone, Michael C
REGISTRATION NUMBER: 39,132
REFERENCE/DOCKET NUMBER: PF
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
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                                           NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS
                                                                        TITLE OF INVENTION:
                                                                                                      APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                         MEDIAL
LIBRARY: PAGE
TONE: 2278458
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: UFILING DATE: HEREWITH CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Palo Alto
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET:
STREET: 3174 Por CITY: Palo Alto
         ADDRESSEE: Incyte rna...acc
                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 94304
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                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
                                                                                                                                                                                                                                                     479
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                                                                                                                                                 INFORMATION:
                                                                                                                                                                                                                                                                                                              l Similarity 100.0%;
13; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                nucleic acid
                                                                                                                                                                                                                                                     GGCTGGTGTCACC 491
                                                                                                                                                                                                                                                                                   GGCTGGTGTCACC 13
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                                                                                                                                                                           Application US/09565910
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3174 Porter Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1742 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Baughn, Mariah R.
WENTION: DELTA 1-PYRROLINE-5-CARBOXYLATE REDUCTASE
                                                                        Baughn, Mariah R.
VENTION: DELTA 1-PYRROLINE-5-CARBOXYLATE REDUCTASE
VENTION: HOMOLOG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          650-855-057
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                                                                                                                  Corley, Neil C.
                                                                                                                                                                                                                                                                                                                                                                                                       PROSNON01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IBM Compatible
                                                                                                                                 fillman,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Incyte Pharmaceuticals, Inc.
                             Incyte Pharmaceuticals,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HOMOLOG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Jennifer L.
                                                                                                                                   Jennifer L.
                                                                                                                                                                                                                                                                                                                            72.2%; Score 13; 100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                            DB 3;
                                                                                                                                                                                                                                                                                                                                            Length 1742;
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COMPUTER READABLE FORM: MEDIUM TYPE: Diskette

94304

USA

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US-08-454-455-3/c
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
            CLASSIFICATION: 530
PRIOR APPLICATION NUMBER: 08/193989
FILING DATE: 09-FEB-1994
PRIOR APPLICATION NUMBER: 08/004142
APPLICATION NUMBER: 08/004142
                                                                                                                                           OPERATING SYSTEM: PC-DOS/MS-DO SOFTWARE: WinPatin (Genentech) CURRENT APPLICATION DATA:
                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 720 Kb floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION: TELEPHONE: 650-855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                  ORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: NOVEL BETA INTEGRIN SUBUNIT
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LIBRARY: PAUL
ONE: 2278458
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Cerrone, Michael C
REGISTRATION NUMBER: 39,132
REFERENCE/DOCKET NUMBER: PF
                                                                                                                                                                                                                                                                                              STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
                                                                                                               FILING DATE:
                                                                                                                              APPLICATION NUMBER: US/08/454,455
                                                                                                                                                                                                                                                                                      STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/09/565,910
                                                                                                                                                                                                                                                   CCUNTRY: UZIP: 94080
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       479 GGCTGGTGTCACC 491
                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 GGCTGGTGTCACC 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity 100.0%; 13; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nucleic acid
DEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Application US/08454455
                                                                                                                                                                                                                                                                                  California
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                                                                                                                                                                                                                                                                   USA
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                                                                                                                                                                                                                                                                                                                                     Genentech,
                                                                                                               30-May-1995
13-JAN-1993
                                                                                                                                                                                                                                                                                                                                                                                                     John W.
                                                                                                                                                                                 PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        72.2%;
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TOPOLOGY:
US-08-454-455-3
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Best Local Similarity
                                 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/67
                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech) CURRENT APPLICATION DATA:
                                                                            REGISTRATION NUMBER: 00,000
REFERENCE/DOCKET NUMBER: PO
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-1994
                                                                                                                                                                                                                                              FILING DATE: 09-FEB-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/004142
                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 720 Kb floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS: ADDRESSEE: Genentech,
                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: McLean, John W. TITLE OF INVENTION: NOVEL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
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                  SEQUENCE CHARACTERISTICS
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                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: FILING DATE: 30-May CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 460 Point San Bruno Blvd CITY: South San Francisco
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                                                  TELEX:
                                                                TELEFAX:
                                                                                                                                                                                                                                             APPLICATION NUMBER:
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TELEFAX: 415/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1012 CTGGTGTCACCTG 1000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Application US/08454455
                                                  910/371-7168
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3789 base pairs
                                                                  415/952-9881
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                                                                                                                                                Wendy M.
                                                                                                                                                                                                                                                                                                                                               30-May-1995
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                                                                                                                 P0699C2D2
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51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 2972;
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RESULT 32
US-08-474-068A-1
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Best Local Similarity
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                                                                                                                         Matches
                                                                                                                                        Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/213,361
FILING DATE: 14-MAY-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/607,293
FILING DATE: 30-0CT-1990
ATTORNEY/ACENT INFORMATION:
NAME: Campbell, Cathryn A.
NAME: Campbell, Cathryn A.
                                                                                                                                                                                                                                                                                                              TELEFAX: (619) 535-894
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Campbell and Flores
STREET: 4370 La Jolla Village Drive, Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION: TELEPHONE: (619) 535-9001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FITLE OF INVENTION:
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                                                                                                                                                                                                  NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/0 FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 4370 La
CITY: San Diego
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 31, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                  TOPOLOGY:
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                                                        976 TGGTGTCACCTGT 988
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                                                                                          4 TEGTETCACCTET 16
                                                                                                                      similarity 100.0%; F
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                                                                                                                                                                                                                                                                                               3959 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                      45..2181
                                                                                                                                                                                                                        CDS
                                                                                                                                                                                                                                                   linear
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                                                                                                                                                                                                                                                                                                                                              535-8949
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -Cadherin Adhesion Molecule
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                                                                                                                                        Score 13; DB 1; Length 3959; Pred. No. 51;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
51;
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                                                                                                                                                                                                                                                                                      RESULT 33
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                                                                                                                                                                                                                                  Sequence 1, Application US/08472481 Patent No. 5863804
                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (619) 535-8949 INFORMATION FOR SEQ ID NO:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 08
FILING DATE: 14-MAY-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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                                                                                                                                                 NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
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TELECOMMUNICATION INFORMATION:
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MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 30-OCT-1990
ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY:
LOCATION:
                                                                                                                                                                                TILE OF INVENTION:
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OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: San Diego
STATE: California
                                                                                  CITY: San Diego
STATE: California
                                                                                                                  STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE:
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                                                                 COUNTRY:
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                                                                                                                E: Campbell and Flores
4370 La Jolla Village Drive, Suite 700
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45..2181
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SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                            Barbara
T-Cadherin Adhesion Molecule
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Pred. No. 51;
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OPERATING SYSTEM: PC-DOS/MS-DOS

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Query Match
Best Local Similarity
Matches 13; Conserv
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: Sequence 39, Application PC/TUS9601314
: GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY:
;
LOCATION:
US-08-472-481-1
                                                                  FILING DATE: 30-JAN-96
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/380,167
FILING DATE: 30-JAN-95
ATTORNEY AGENT INFORMATION:
NAME: John W. Freeman
REGISTRATION NUMBER: 29,066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (619) 535-8949
(INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 3959 base pairs
                                                                                                                                                                                                      COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5" Diskette, 1.44 Mb

COMPUTER: IBM PS/2 Model 50Z or 55SX

OPERATING SYSTEM: MS-DOS (Version 5.0

SOFTWARE: WordPerfect (Version 5.1)

CURRENT APPLICATION NUMBER: PCT/US96/01314
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 14-MAY-PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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APPLICATION NUMBER: US 08/213,361
FILING DATE: 14-MAY-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-LJ 1686
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PILING DATE: 30-0CT-1990
ATTORNEY/AGENT INFORMATION:
NAME: Campa-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: METHODS FOR IDENTIFYING INTEGRIN TITLE OF INVENTION: ANTAGONISTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                 REFERENCE/DOCKET NUMBER: 00 
TELECOMMUNICATION INFORMATION: 
TELEPHONE: (617) 542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TUMBER OF SEQUENCES:
TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                            STATE: Massachusetts COUNTRY: U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                 E: Fish & Richardson P.C.
225 Franklin Street
   (617)
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; Sequence 39, Application US/08476062A
; Patent No. 5877275
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Query Match
Best Local Similarity
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                           INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 07-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/2:
FILING DATE: 21-MAR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatil
OPERATING SYSTEM: Wind
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LENGTH: 5137 base pairs
                                                                                        FEATURE:
                                                                                                    MOLECULE TYPE:
                                                                                                                                                                                                                                        TELEPHONE: 617/542-5070
                                                                                                                                                                                                                                                                                                               FILING DATE: 18-JUN-1990
APPLICATION NUMBER: 07/2:
FILING DATE: 28-JUN-1988
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                         NAME/KEY: Coding Sequence LOCATION: 95...3604
                                                                                                                                 STRANDEDNESS:
                                                                                                                                              TYPE: nucleic acid
                                                                                                                                                                                                                           TELEFAX:
                                                                                                                                                                                                                                                                                    NAME: Freeman, John W
REGISTRATION NUMBER: :
                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: FILING DATE: 04-JA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Boston
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                                                                                                                       TOPOLOGY:
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02110-2804
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SYSTEM: Windows95
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SS: single
                                                                                                                       linear
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72.2%;
100.0%;
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CONTROLLING CELLULAR IMMUNE/INFLAMMATORY
RESPONSES WITH BETA2 INTEGRINS
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Score 13; ; Pred. No.
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Pred. No.
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 DB 2;
50;
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50;
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             Length 5138;
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Db 4209 GGCTGGTGTCACC 4197	Qy 1 GGCTGGTGTCACC 13	Matches 13; Conservative 0; Mismatches 0; Indels
		0;
		0; Gaps 0;
		0;
		420

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Result
No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Minimum DB
Maximum DB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Title:
Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Database :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OM nucleic - nucleic search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Word size
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Post-processing: Listing first 120 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Total number of hits satisfying chosen parameters:
                                                                                                                                                           Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                       Score
   1661168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           length: 0
length: 2000000000
                                                                                                                                                                                           N Geneseq 19Jun03:*

1: /SIDS1/gcgdata/geneseq/geneseqn-emb1/NA1980.DAT:*
2: /SIDS1/gcgdata/geneseq/geneseqn-emb1/NA1981.DAT:*
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EBV gp350/220 cDNA
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Murine C16N-2 cDNA
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Murine C16N-1 codin
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Ovary cancer relat
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Probe #380 used to
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AAS72881

AAS32249 AAK71839

Human DNA repair a

immune/haema

transporter

Novel Human DNA encoding DNA encoding novel gene express

human DNA re

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AAQ91011
XX AAQ91011
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XX Primer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9516028-A1.
                               Compsn. useful for polynucleotide synthesis and cyclical amplification - comprising a mixt. contg. an enzyme with exo-nuclease activity and a DNA polymerase with less 3'-5' exo-nuclease activity than the enzyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         hybridizes to Epstein-Barr virus (EBV) nucleic acid, for detecting EBV in clinical specimens to determine patients at high risk of to developing EBV infections -
                                                                                                                                                                                  WPI; 1995-224316/29.
                                                                                                                                                                                                                                             Mullinax RL, Sorge JA;
                                                                                                                                                                                                                                                                                                                                                                 16-FEB-1994;
08-DEC-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-DEC-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Primer; PCR; amplification; DNA polymerase; exonuclease; Pfu; Taq; Klenow fragment; T4; T7; Deep Vent; synthesis; mismatch; human; antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Primer binding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; Page 44; 59pp; English.
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hain variable region; ss.
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Pred. No. 0.41;
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Examples;

Page 35;

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English

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A DNA protein-binding assay is provided, useful for screening libraries of synthetic or biological cpds. for their ability to bind DNA test sequences. The assay is versatile in that any number of test sequences can be tested by placing the test sequence adjacent to a defined protein-binding screening sequence. Binding of mols. to these test sequences changes the binding characteristics of the protein mol. to its cognate binding sequence. When such a mol. binds the test sequence, the equilibrium of the DNA:protein complexes is disturbed, generating changes in the concentration of free DNA prob
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This primer binds to a region at the 5' end of the Epstein-Barr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA protein-binding assay; test sequence; screening sequence, promoter; target; TATA box; Herpes Simplex Virus; HSV; origin of replication; UL9; transcription factor; TFIID: ds.
                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence-directed DNA-binding molecules
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-DEC-1992;
17-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8
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93US-0123936.
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Best Local S
Matches 16
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27-JUN-1991;
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17-SEP-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Duplex TFIID;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAT64258 standard; DNA; 50
                                                                                                                                                                                                                                                                                              Altering binding characteristics of DNA binding proteins to duplex DNA - by attaching specific small cpd. to target region close to the protein's binding site, useful in treatment of viral disease, cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cancer; inherited genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EBV L2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-MAR-2003
17-MAR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT64258;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             One application of this method is to eucaryotic general transcription
                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1997-020402/02.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Andrews
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-DEC-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (GENE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (Updated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA; target region; binding characteristic; DNA binding protein; transcription factor; binding site; inhibition; enhance; inherited genetic disorder; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENELABS TECHNOLOGIES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        ВМ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GGCTGGTGTCACCTGT
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     site 90021) TFIID binding site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              92US-0996783
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               93US-0171389
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 16; pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10 G;
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as target regions in the method of the invention. The method for altering the binding protein to DNA comprises contacting the duplex DNA with a small molecule whi binds sequence-specifically to a target region, where, when the s molecule is bound to the target region, it is adjacent to, but no overlapping by more than 4 bp, a binding site for a DNA-binding or the small molecule is added at a concentration effective to alter

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The sequences given in AAT63713-4312 represent

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27-JUN-1991;
23-DEC-1992;
17-SEP-1993;
07-JUN-1995;
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7546/c
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Sequences AAX17001 to AAX17600 represent specifically claimed target test sequences that are used in the method of the invention of determining the DNA sequence preference of DNA-binding molecule. The method comprises: (i) adding a test molecule and a DNA-binding protein to a mixture of duplex DNA test oligonucleotides, each of the test oligonucleotides, having a test sequence adjacent to a screening sequence, where the screening sequence binds to the DNA-binding protein with a binding affinity that is independent of the DNA sequence of the test
                                                                                                                                                                                                                                                               Claim
                                                                                                                                                                                                                                                                                                                    sequence attached
                                                                                                                                                                                                                                                                                                                                            Determination of DNA sequence preference of based on inhibition of binding of protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-JUN-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Test sequence; DNA-binding molecule; screening sequence; human; nucleic acid amplification; target; viral; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Andrews
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                                                                                                                                                                                                                                                               Columns 379-380;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cantor CR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               from Epstein Barr virus L2 (start site 90021).
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92US-0396783.
93US-0123936.
95US-0475228.
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-DEC-1993;
07-JUN-1995;
27-JUN-1991;
23-DEC-1992;
17-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA binding molecule screening; inhibition of transcription; infection; human immunodeficiency virus; HIV; parasite; cancer; cardiovascular; respiratory; gastrointestinal; endocrine; metaborheumatic; immunological; haematological; neurological; psychiatric; dermatological; ophthalmological; musculo-skeletal;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-JUL-1999;
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Pred. No.
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The present sequence is one of a large number of 5' ESTs derived from mRNAs encoding secreted proteins. An ORF has been identified within the sequence. The 5' ESTs were prepared from total human RNAs or polyA+ RNAs derived from 30 different tissues. EST sequences usually correspond mainly to the 3' untranslated region (UTR) of the mRNA because they are often obtained from oligo-dT primed cDNA illoraries. Such ESTs are not well suited for isolating cDNA sequences derived from the 5' ends of mRNAs and even in those cases where longer cDNA sequences have been obtained, the full 5' UTR is rarely included. 5' ESTs are derived from mRNAs and genomic DNAs. 5' ESTs are also used to obtain full length cDNAs and genomic DNAs. 5' ESTs are also used in diagnostic, forensic, gene therapy and chromosome mapping procedures. They are used to obtain upstream regulatory sequences and to design expression and secretion vocators.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and fo diagnostic, forensic, gene therapy and chromosome mapping procedu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human secreted protein 5'
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P-PSDB; AAG01770.
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1;
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Pred. No.
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Sequence 455 BP;

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Best Local Similarity
Matches 16; Conserv
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21-JUN-2000; 2000US-213361P.
01-SEP-2000; 2000US-229763P.
05-SEP-2000; 2000US-230629P.
14-SEP-2000; 2000US-232565P.
19-DEC-2000; 2000US-27037P.
08-JAN-2001; 2001US-260796P.
                                                                             determining cancer in a patient. (I), (II) or antigen-presenting cells expressing (II) is useful for stimulating and/or expanding T cells specific for a tumour protein. The method comprises contacting T cells with one of the components under conditions to permit the stimulation and/or expansion of the cells. A composition comprising (I) is useful for stimulating an immune response in a patient and for inhibiting the development of a cancer especially lung cancer in a patient. An isolated T cell population is useful for removing tumour cells from the biological sample and for inhibiting the development of cancer in a patient. AAS61874 represent novel human lung small cell
                                                                                                                                                                                                                                                                                                                                                                      polynucleotides (I) and polypeptides (II) used in a method of detection of a patient. The method is optionally performed by utilising oligonucleotides (III), where the biological sample from the patient is contacted with (III), detecting the amount of polynucleotide hybridised to (III) in the sample and comparing the amount of polynucleotide to a predetermined cut-off value and there determining cancer in a nation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 435 BP; 105 A; 114 C; 133 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lung small cell carcinoma antigen, cDNA #63.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAS61522;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lung tumour polynucleotide and polypeptides useful in therapy and diagnosis of cancer especially lung cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-010896/01.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-JAN-2002
                                                             cancer antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to novel isolated lung small cell cancer antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 146; 295pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (CORI-) CORIXA CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Wang T,
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Pred. No.
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                                                                of the invention.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to human polynucleotides (AAI79941-AAI93841) and the encoded proteins (AA000010-AA013910) that exhibit activity elating to cytokine, cell proliferation or cell differentiation or which may induce production of other cytokines in other cell populations. The polynucleotides and polypeptides are useful in gene therapy, vaccines or peptide therapy. The polypeptides have various cytokine-like activities, e.g. stem cell growth factor activity, haematopolesis regulating activity, tissue growth factor activity, immunomodulatory activity and activity inhibin activity and may be useful in the diagnosis and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Isolated nucleic acids and polypeptides, useful for preventing diagnosing and treating e.g. leukaemia, inflammation and immundisorders -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; cytokine; cell proliferation; cell differentiation; gene therapy; vaccine; peptide therapy; stem cell growth factor; haematopoiesis; tissue growth factor; immunomodulatory; cancer; leukaemia;
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                                                                                                                                                                                                                                            Sequence 675
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; SEQ ID NO 9164; 1399pp + Sequence Listing; English
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18-MAY-2000; 2000US-0577409.
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                                                                                                                                                                                                                                                                                                                                     Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-FEB-2001; 2001WO-US04927.
                                                                                                                                                                                                                                                                                                                                                                                                                             treatment of cancer, leukaemia, nervous system disorders, arthritis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                  ftp.wipo.int/pub/published_pct_sequences.
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CTGGTGTCACCTGTTA
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                                                                                                                                                                                                                                            BP; 189 A; 165 C; 147 G; 172 T; 2 other;
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                                                                                                                                                 Score 16; DB 22;
Pred. No. 5.5;
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This DNA encodes a human secreted protein. The specification provides correted protein sequences (AAW3363) to AAW4369) encoded by the nucleic cacid sequences shown in AAV43611 to AAV43619. The invention-provides a cumethod of identifying a secreted polypeptide which is modified by rough committees. The secreted proteins can be used in assays to determine the biological activities, such as cytokine, cell proliferation, or cellular ciphibin activity, chemotactic or chemokinetic activity, haemostatic or chrombolytic activity, receptor/ligand activity, tumour inhibition, or cellular compounds which inflammatory activity. The proteins can also be used as corrected proteins, compounds which affect the biological activities of the groteins. Compounds which affect the biological activities of the corrected proteins or their ability to interact with specific ligands can be used proteins or their ability to interact with specific ligands can compound to the proteins in screening assays. The proteins and compounds which affect the biological activities of the proteins and compounds which affect the biological activities of the compounds which affect the biological activities of the compounds which affect the biological activities of the proteins and compounds which affect the biological activities of the com
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Secreted protein; human; cell proliferation; cytokine activity; tissue growth; cellular differentiation; regeneration; activin; inhibin; chemotactic; haemostatic; thrombolytic; tumour inhibit anti-inflammatory activity; biomarker; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Secreted human polypeptides - having cytokine, cell proliferation differentiation, activin or inhibin, tumour inhibition or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-DEC-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAV43611 standard; DNA; 1142 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 6; Page 38; 78pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Escobedo J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-JUN-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /product= "human secreted protein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Williams
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Best Loc Matches Query Match Best Local (

Similarity

88.9%; Score 16; 100.0%; Pred. No.

DB 19; 5.5;

Length 1142; Indels

Conservative

<u>.</u>

Mismatches

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Gaps

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Sequence 1142 BP; 304 A; 335 C;

331 G;

172 T; 0 other;

밁

159

0

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AAV62754/c
ID AAV62754 standard; cDNA; 1152
                           It shows some similarity to database sequences. The invention provides polynucleotides (see AAV62746-55) from human foetal brain, adult testis, adult brain, adult kidney and feetal kidney (all deposited as composite clone ATCC 98404), which encode human secreted proteins (see AAW74720-29). The polynucleotides and proteins are predicted to have biological activities which would make them suitable for treating, preventing or ameliorating medical conditions in humans and animals, although no supporting data are given. Suggested activities include nutritional activity, immune stimulating (e.g. as vaccines) or suppressing activity, activity, activity, chemotactic/chemokinetic activity, activity, cativity, and tumour inhibition activity. The polynucleotides are also stated to be useful for gene therapy, and can be used in recombinant production of the polypeptidess.
                                                                                                                                                                                                                                                                                                                                Full-length cDNA clone fm150_1 includes an open reading frame encoding a human secreted protein (see AAW74728). It was isolated from a human adult brain cDNA library using methods which are selective for cDNAs encoding secreted proteins, or was identified as encoding a secreted or transmembrane protein on the basis of as encoding a secreted or transmembrane protein on the basis of computer analysis of the amino acid sequence of the encoded protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 34; Page 95; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New polynucleotide(s) encoding secreted human proteins - are derived from, e.g. human foetal brain or foetal kidney cDNA libraries, potentially useful as, e.g. vaccines or thrombolytic agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13-APR-1998;
15-APR-1997;
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Sequence 1152 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Racie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Agostino MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GEMY ) GENETICS INST INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Spaulding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Jacobs
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97US-0843374.
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11..973
/*tag= a
342 A; 329 C; 317 G; 164 T; 0
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Treacy M;
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 other;
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Query Match

88.9%;

Score 16;

DB 19;

Length 1152;

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RESULT 12
ABQ92057/c
ID ABQ920
XX Human;
XX Human;
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XX Im
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to human secreted or transmembrane protein (I), their fragments and is encoded by specific complementary deoxyribonucleic acid (cDNA) inserts (II), where the protein is substantially free from other mammalian proteins. (I) are useful for preventing, treating or ameliorating a medical condition, especially immunological treatment or prevention of tumours. (I) exhibits activity relating to angiogenesis, cytokine, cell proliferation, cell differentiation, antiinflammatory, stem cell growth factor activity and activin or inhibin-related activities. (I) can be used to manipulate stem cells in culture to give rise to neuroepithelial cells that can be used to augment or replace cells damaged by illness, autoimmune disease, accidental damage or genetic disorders. (I) induces the proliferation of neural cells and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Jacobs K,
Merberg D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABQ92057 standard; cDNA; 1152
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                                                                                                                                                                                                                                                                                                                                             Claim 189; Page 195-196; 284pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                            protein, useful for diagnosis and treatment cancer, autoimmune diseases, bone disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel secreted or transmembrane protein and protein, useful for diagnosis and treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    P-PSDB; ABP61843.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EVAN/)
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MERBERG D.

TREACY M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COLLING-RACIE
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LAVALLIE E R.
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, Treacy M,
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Spaulding V;
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                                                                                                                                                                                                                                                                                                                                                                                                                         polynucleotide encoding the of neurological disorders, and lung or liver fibrosis
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RESULT 13
AAC77733/G
ID AAC77733/G
AC AAC77
XX AAC77
XX AAC77
XX AAC77
XX Human
XX Human
XX Human
XX Human
XX diagn
KW diagn
KW antii
KW vasot
KW inmur
KW allen
KW haelen
KW haelen
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Best Local S
Matches 16
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AAC77733 standard; cDNA; 1217 BP
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   Claim 1; Page 709; 2352pp; English
                                                                Novel isolated nucleic acids comprising sequences encoding peptides useful for treating or diagnosing e.g.\ 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   diagnosis; cytostatic; proliferative; vulnerary; immunomodulator; antityiral; antitheteric; antiarthritic; antityiral; antitheteric; antityiral; antitheteric; antibacterial; cardiant; antithinflammatory; antithyroid; antiallergic; antibacterial; cardiant; dermatological; neuroprotective; thrombolytic; coagulant; nootropic; vasotropic; antipsoriatic; antiangiogenic; gene therapy; inflammation; immune disorder; haematopoietic cell disorder; autoimmune disorder; allergic reaction; graft versus host disease; organ rejection;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human cancer associated gene sequence SEQ ID NO:127.
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                                                                                                                                                                                                                                                                                              Rosen CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-MAR-2000; 2000WO-US05882
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              haemostatic; thrombolytic; cardiovascular disorder; infection; neurological disease; drug screening; ss.
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                                                                                                                                                                                                                                                                                                                                                                          (HUMA-) HUMAN GENOME
                                                                                                                                                                                   2000-587533/55.
DB; AAB43524.
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                                                                                                                                                                                                                                                                                              Ruben SM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   99US-0124270.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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RESULT 14
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Best Local S
Matches 16
                                                                                    Jones WL, Kassa
Lamson G, Leshk
Stache-Crain B,
                                                                                      Crkvenjakov R, Dickson M, Drmanac N, Escobedo J, Garcia PD, Garcia V, Giese K, Escobedo J, Kassam A, Kennedy GC, Kita D, La Jones WL, Kassam A, Kennedy GC, Kandazzo F, Williams
                                                                                                                                                                                                                                                                                                                                                                       28-JAN-1998;
24-FEB-1998;
31-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Polymicleotides, polypeptides, antibodies, agonists and antagonists from the present invention may be used to treat immune disorders by activating or inhibiting the proliferation, differentiation or mobilisation of immune cells, to treat disorders of haematopoietic cells, autoimmune disorders, allergic reactions, graft versus host disease and organ rejection, modulate haemostatic or thrombolytic activity, modulate inflammation, cancers, cardiovascular disorders, neurological disease and bacterial or viral infections. The peptides, nucleotides, antibodies, agonists and antagonists may be also be used in drug screens. AAC78449 to AAC78457 and AAB44240 represent sequences used in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; gene; gene expression product; diagnosis; therapy; probe; detection; mapping; tissue typing; profiling; forensic; cancer; genetic analysis; colorectal cancer; breast cancer; lung cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human gene expression product cDNA sequence SEQ ID NO:5243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           dermatological, neuroprotective, cardiant, thrombolytic, coagulant, nootropic, vasotropic, antipsoriatic and antiangiogenic. The polynucleotides and polypeptides can be used for preventing, treating ameliorating medical conditions and diagnosing pathological conditions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antidiabetic; antiasthmatic; antirheumatic; antiarthritic;
antiinflammatory; antithyroid; antiallergic; antibacterial; antiviral;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAC77607 to AAC78448 encode the human cancer associated proteins git
in AAB43398 to AAB44239. The proteins can have activities based on t
tissues and cells the genes are expressed in Example of activities
include: cytostatic; proliferative; vulnerary; immunomodulator;
                                  WPI; 1999-494092/41
                                                                                                                                                                                                                                                         (CHIR )
(HYSE-)
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                                                                                                                                                                                                                                                         HYSEQ INC.
                                                                                                                                                                                                                                                                                   CHIRON CORP.
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                                                                                                                                                                                                                                                                                                                                                                    98US-0080666.
98US-0072910.
98US-0075954.
98US-0080114.
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                                                                                                                                                                                                  Drmanac
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Labat I;
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                                                                                                                      Reinhard
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Novel human genes and their expression products which differentially expressed in different cell types

2479pp;

English

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RESULT 15
ACC46555/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
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            28-MAR-2001;
29-MAR-2001;
29-MAR-2001;
16-MAY-2001;
17-MAY-2001;
17-MAY-2001;
19-JUN-2001;
                                                                                                                                                                                                                                                                                          connective tissue disorder; drug screening; proteome analysis; gene therapy; antisense therapy; genotyping; transgenic animal; disease model; toxicological testing; transcript imaging; secreted protein; extracellular matrix; gene; ss.
                                                                                                                                                                                                                                                                                                                                                            Human; dithp; diagnostic and therapeutic polynucleotide; diagnosis; cancer; cell proliferative disorder; autoimmune disorder; inflammatory disorder; infection; hormonal disorder; metabolic disorder; neurological disorder; gastrointestinal disorder; transport disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ACC46555 standard; cDNA; 2382
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 1278 BP;
                                                                                                                                                      27-MAR-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human dithp
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                                             2001US-280067P.
2001US-280068P.
2001US-291280P.
2001US-291829P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      secreted/extracellular matrix protein-encoding
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            2001US-291849P.
2001US-299428P.
                                                                                                                                                        2002WO-US10056
                                                                                                                     2001US-279619P
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100.0%;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              282 T; 10 other;
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5.5;
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DNA sequence

encoding Epstein-Barr

virus

(EBV)

outer surface

protein

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AAN50114/c
ID AAN501
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AC AAN501
XX
DT 25-MAR
DT 17-OCT
XX
DE DNA se
XX
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                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to novel human diagnostic and therapeutic CC polynucleotides designated dithp (ACC46080-ACC46749) and to their CC encoded proteins (DITHP; ABR41136-ABR41812). The invention also relates CC to polynucleotide sequences at least 90% identical to the dithp cDNA CC sequences of the invention; recombinant vectors, host cells and CC transgenic organisms comprising a dithp nucleic acid sequence; the CC recombinant production of DITHP proteins; antibodies specific for DITHP CC proteins; microarrays comprising dithp nucleic acid sequences; methods of screening CC of detecting dithp nucleotide and protein sequences; methods of screening CC for compounds which specifically bind a DITHP protein; and methods of a wide variety of conditions including cancer and other cell compressing the toxicity of test compounds using a dithp hybridisation for probe. Dithp nucleic acid sequences and DITHP proteins may be used in the CC diagnosis of a wide variety of conditions including cancer and other cell compounds using a dithp hybridisation of the creation of protein activity or gene expression. DITHP proteins may be used in the CC disorders; neurological disorders; patternintestinal disorders; transport CC disorders; and connective tissue disorders. They may also be used to screen for modulators of protein activity or gene expression. DITHP proteins can additionally be used in analysis of the proteome of a tissue or cell type and to induce antibodies. The dithp nucleic acids are additionally useful in somatic or germline gene therapy of the disorders can be used to the generation of transgenic animal models of human disease or knock in the generation of transgenic animal models of human disease or knock in the present sequence data for this patent did not form part of the printed cartification, but was obtained in electronic format directly from WIPO art fer winc it from WIPO
                                                                                                                                                                       RESULT 16
                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 16
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                                                                                                                                                                                                                                                                                                                                                          Sequence 2382 BP; 456 A; 1032 C; 337 G;
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                                                25-MAR-2003
17-OCT-1991
                                                                                                                                     AAN50114 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim
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                                                                                                                                                                                                                                                                                                                                                                                          ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                       1869
                                                                                                                                                                                                                                          1 GGCTGGTGTCACCTGT 16
                                                                                                                                                                                                                                                                                         16;
                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABR41618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID No 476;
                                                                                                                                                                                                                       GGCTGGTGTCACCTGT 1854
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hillman JL,
                                                                                                                                                                                                                                                                                          Conservative
                                                (updated)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Œ,
                                                                                                                                     DNA;
                                                                                                                                                                                                                                                                                                       88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tran AB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ran AB, Dahl CR, Giet,
, Yu JY, Tuason O, Yo
Liu TF, Nguyen DA,
Lewis SA, Chen AJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lo A,
                                                                                                                                     2721
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  591pp; English.
                                                                                                                                                                                                                                                                                                       Score 16; Pred. No.
                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lan RY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Urashka ME;
                                                                                                                                                                                                                                                                                                         DB 25;
5.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gietzen D, C
O, Yap PE, Aı
DA, Kleefeld
                                                                                                                                                                                                                                                                                                                                                          556
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              y binds to a polypeptide polynucleotide, and to
                                                                                                                                                                                                                                                                                                                                                          H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Panzer
                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                            1 other;
                                                                                                                                                                                                                                                                                                                       Length 2382;
                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Chinn J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Amshey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gerstin
                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SR;
                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ø
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RESULT 17
AATO4821/c
ID AATO4
XX AATO4
XX AATO4
XX IB-JA
DT 18-JA
DT 18-JA
DX EBV g
XX EBV;
XX EBV;
XX EBV;
YX EBV;
YX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      á
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                     misc_feature
                                                                                                                                                                mat_peptide
                                                                                                                                                                                                        sig_peptide
                                                                                                                                                                                                                                                                                                                    Epstein-Barr virus
                                                                                                                                                                                                                                                                                                                                                                                                        EBV gp350/220 cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                     18-JAN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAT04821;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAT04821
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 2721 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The sequence encodes an outer surface viral protein of EBV, to generate antibodies reacting with the surface proteins of EBV-infected cells, and in the preparation of a vaccine agai (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 21-23; 26pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New fragment of Epstein-Barr Virus DNA - useful in vector to express polypeptide for use in prepn. of vaccine against the virus and for use in diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-JUL-1984;
30-JAN-1984;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kleff E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EP151079-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    mat_peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Epstein-Barr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (UYCH-) UNIV CHICAGO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-JAN-1985;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Epstein-Barr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1985-191978/32.
DB; AAP50073.
                                                                                                                                                                                                                                                                                                                                                             gp350; gp220; gp350/gp220; non-splicing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GGCTGGTGTCACCTGT 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GGCTGGTGTCACCTGT 2109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tanner J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          virus; antigen; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              84US-0633558.
84US-0575352.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                85EP-0400141
/*tag= d
/function= splice donor site
/function=splice donor site
/note= "bases 2513-2517 (AAGT) are replaced
/note= "GTCA in the non-splicing variant"
                                                                                                                  /*tag= b
1068..3734
/*tag= c
2514..2515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /*tag=
/label=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
1..2721
                                                                                                                                                                                                          /*tag= a
1014..1067
                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         762 A; 876 C; 557 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CDNA; 5931 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          88.9%; 50
100.0%; Pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                     entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hummel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          a
EBV surface
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Σ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 16;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Belsel C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         526 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 6
5.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              6;
                                                                                                                                                                                                                                                                                                                                                               variant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ٥.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 2721;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                 vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      against EBV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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RESULT 18
ABA15144
ID ABA15
XX ABA151
XX ABA15
XX ABA15
XX ABA15
XX BLuma
XX Huma
XX Huma
XX Huma
XX Huma
XX Huma
XX Immu
XX ant:
XW AN:
XW AN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The donor and acceptor splice sites of the EBV gene encoding gp350/220 are mutated by replacement of nature uncleotides by non-native nucleotides, without altering the encoded amino acid sequence, resulting in elimination of gp220 prodn. Recombinant homogeneous gp350, useful in vaccines, is expressed in mammalian or insect cell
                                                                                                                                                                                                                                                       immunosuppressive; antiinflammatory; anti-HIV; antibacterial; vulnerary; antiparkinsonian; antisikling; antianaemic; antiarthritic; cancer; antirheumatic; hepatotropic; cerebroprotective; antiinflammatory; antiallergic; antidiabetic; antiulcer; anticonvulsant; antifungal; antiparasitic; cardiant; immune disorder; cardiovascular disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 5931 BP; 1453 A; 1782 C; 1437
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New DNA encoding a homogeneous gp350 protein - can preventing and treating Epstein-Barr virus-related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           misc_feature
                                                                                                                                                                                                                                neurological disease; infection;
                                                                                                                                                                                                                                                                                                                                                                                    Human; nootropic; neuroprotective; cytostatic; dermatological; virucide;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Human nervous system related polynucleotide SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABA15144;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABA15144
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 2; Fig.1; 61pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Jackman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18-APR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9528488-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (AVIR-) AVIRON.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1995-373802/48.
DB; AAR80144.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GGCTGGTGTCACCTGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GGCTGGTGTCACCTGT 3122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3105..3106

/*tag= e
/function= splice acceptor site
/note= "bases 3104-3107 (AGGT) are replaced?
/note= "bases 3104-3107 (AGGT) are replaced?
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              94US-0229291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             95WO-US04611.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA; 13559
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                  nephrotropic; gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   G; 1259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 5931;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         be used f
                                                                                                                                                                                                                                therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                        7475
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         e or
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17-JAN-2001; 2001WO-US01334.

16-AUG-2001.

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22-AUG-2000

23-AUG-2000

30-AUG-2000

01-SEP-2000

01-SEP-2000

01-SEP-2000

05-SEP-2000

06-SEP-2000

08-SEP-2000

08-SEP-2000

08-SEP-2000

08-SEP-2000

11-SEP-2000

21-SEP-2000

21-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18-AUG
 2000US-0189874

2000US-0199173

2000US-0199173

2000US-0209467

2000US-0214886

2000US-0214886

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2000US-0214896

2000US-0218296

2000US-0224519

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2000US-0223447

2000US-0231444

2000US-0231468

2000US-0231474

2000US-0231498

2000US-02334974

2000US-023498

2000US-0234894

2000US-0234898

2000US-0234998

2000US-0234998
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2000US-0180628.
2000US-0184664.
2000US-0186350.
   17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
01-DEC-2000;
01-DEC-2000;
05-DEC-2000;
05-DEC-2000;
06-DEC-2000;
08-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-OCT-2000;
20-CCT-2000;
01-NOV-2000;
08-NOV-2000;
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02-0CT-2000

02-0CT-2000

02-0CT-2000

13-0CT-2000

13-0CT-2000

20-0CT-2000

20-0CT-2000

20-0CT-2000

20-0CT-2000
Nucleic acids encoding 3224 human nervous system antigen polypeptides, useful for preventing, diagnosing and/or treating nervous system cancers and metastases -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             L7-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          7-NOV-2000
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                                                                      2001-541565/60.
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2000US-0246476.
2000US-0246478.
2000US-0246523.
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2000US-0246527.
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2000US-0246527.
2000US-0246610.
2000US-0246610.
2000US-02466113.
                                                                                                       Barash SC,
                                                                                                                                                                          2000US-0251391
2000US-02513030
2000US-02513030
2000US-02513988
2000US-0251479
2000US-0251479
2000US-0251856
2000US-0251868
2000US-0251869
2000US-0251990
2000US-0251990
2000US-0251997
2000US-0254097
2001US-0259678
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2000US-0249209

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2000US-0249211

2000US-0249211

2000US-0249213

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2000US-0249214

2000US-0249215

2000US-0249216

2000US-0249217

2000US-0249217

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2000US-0249295
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2000US-0237049.
2000US-0237949.
2000US-0239935.
2000US-0239937.
2000US-0241785.
2000US-0241786.
2000US-0241787.
2000US-0241808.
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2000US-0246474
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2000US-0242221
                                                                                                                                          GENOME
                                                                                                                                          SCI
                                                                                                         Ruben
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Disclosure; SEQ

IJ

NO 7475; 1701pp + Sequence Listing; English.

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RESULT 19
ABN52794
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Best Lccal S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The nucleic acids, proteins, antibodies and (ant)agonists are useful in the diagnosis, treatment and prevention of: (a) cancer, e.g. breast and ovarian cancer and other cancers of the adrenal gland, bone, bone marrow, breast gastrointestinal tract, liver, lung, or urogenital; (b) immune disorders e.g. Addison's disease, allergies, autoimmune haemolytic anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's disease, multiple sclerosis, heumatoid arthritis and ulcerative colitis; (c) cardiovascular disorders such as myocardial ischaemias; (d) wound healing; (e) neurological diseases e.g. cerebral anoxia and epilepsy; and (f) infectious diseases such as viral, bacterial, fungal
                                                                                            New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription a genome, useful for detecting tissue-, pathology-, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ABB14678-ABB18001) useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. The genes are isolated from a range of human tissues disclosed in the specification.
                                                                                                                                                                                                                                               28-JUL-2000; 2000US-221607P.
02-MAY-2001; 2001US-287724P.
                                                                                                                                                                                                                                                                                                                                                                                                                           Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mouse spliced transcript detection oligonucleotide SEQ ID NO:25542
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 13559 BP; 2882 A; 3867 C; 4239 G;
                                                                                  developmental-specific genes
                                                                                                                                                           WPI; 2002-257383/30
                                                                                                                                                                                      Shoshan
                                                                                                                                                                                                                                                                                          20-JUL-2001; 2001WO-IB01903
                                                                                                                                                                                                                                                                                                                         07-FEB-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABN52794;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABN52794 standard; DNA; 65
                                                                                                                                                                                                                   (COMP-) COMPUGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12761
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              parasitic infections.

e: The sequence data for this patent did not form part of the nted specification, but was obtained in electronic format directions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16;
                                                                                                                                                                                                                                                                                                                                                                                                               variant;
                                                                                                                                                                                                                                                                                                                                                                                                                           mouse; rat; splice transcript; detection; RNA transcript;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GGCTGGTGTCACCTGT
                                                    SEQ ID 25542; 47pp;
                                                                                                                                                                                      Wasserman A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                               transcriptome;
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Pred. No.
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The present invention describes oligonucleotide libraries for detecting messenger RNAs that populate a (sub-)transcriptome, where the (sub-)transcriptome comprises messenger RNAs transcribed from multiple

Claim 1; Page 10906; 11750pp; English

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RESULT 20
ABV56544
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Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 quantitatively characterising the corresponding transcriptome, and in detecting RNA transcripts and splice variants of human or animal transcriptomes. The libraries may also be used as specialised mini libraries to detect transcripts of a sub-transcriptome under a particular biological or pathological state, and so allowing the detection of tissue—and pathology-specific genes such as those genes only expressed in specific tissue under a specific pathological condition; to detect developmental specific genes; and to detect RNA transcripts and splice variants of a transcriptome of a patient suffering from a particular disorder. ABNZ-7253 to ABNS-9589 represent chief exemplification of the present invention.

N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                           17-FEB-2000; 2000US-183319P.
16-MAR-2000; 2000US-189862P.
25-MAY-2000; 2000US-207454P.
09-JUN-2000; 2000US-211314P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the genome, which encodes one or more messenger RNA splice variants. The oligonucleotide libraries are useful for detecting mRNAs from a biological sample, in expression profiling studies, in qualitatively or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            transcription units that populate a genome. The library comprises several oligonucleotides, each capable of hybridising selectively set of messenger RNAs transcribed from a given transcription unit
                                         Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, usef for detecting presence of prostate cancer
                                                                                                       WPI; 2001-662795/76
                                                                                                                                    Schlegel R,
                                                                                                                                                                                                                                                                                                         20-FEB-2001; 2001WO-US05171
                                                                                                                                                                                                                                                                                                                                        23-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                   WO200160860-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                              pharmacogenomic
                                                                                                                                                                                                                                                                                                                                                                                                                                               Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human prostate expression marker cDNA 56535.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABV56544;
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                                                                                                                                                                  (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                              prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15;
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2000US-211314P.
2000US-219007P.
2000US-255281P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                 marker;
                                                                                                                                                                                                                                                                                                                                                                                                                                              cancer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                 gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                              cytostatic; carcinogen;
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                                                                                                                                     Monahan JE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23 G; 23 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           <u>,,</u>
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                                                                                                                                                                                                                                                                                                                                                                                                                                               pharmacodyanamic marker;
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ARESULT 21
AAF11311
ID AAF11311
AC AAF11
AC AAF1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Multiple gene expression; filamentous fungal cell; EST; expressed sequence tag; Fusarium venenatum; Aspergillus niger; Aspergillus oryzae; Trichoderma reesei; identification; recombination; culture condition; environmental stress; spore morphogenesis; metabolic pathway engineering; catabolic pathway engineering; ss.
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(d) assessing the efficacy of a therapy for inhibiting prostate cancer in a patient;
(e) selecting a composition for inhibiting prostate cancer in a patient (f) assessing the prostate cell carcinogenic potential of a compound;
(g) determining whether prostate cancer has metastasized in a patient;
(h) assessing the aggressiveness or indolence of prostate cancer in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to an isolated nucleic acid molecule (I) comprise a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the specification or its complement. (I) is useful for:

(a) assessing whether a partient is affiliated with prostate cancer;

(b) monitoring the progression of prostate cancer in a patient;

(c) assessing the efficacy of a test compound to inhibit prostate cancer in a patient;
                    The present invention describes a method for monitoring differential expression of genes in a first filamentous fungal (FP) cell relative expression of the same genes in one or more second filamentous fungal cells. The method uses fluorescence-labeled nucleic acids isolated fithe FF cells and a substrate of expressed sequence tags (EST). The Exame used in the methods for monitoring differential expression of general uses in the methods for monitoring differential expression of general sequence.
                                                                                                                                                                                                                                                                                                      Monitoring differential expression of genes in fi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                          Claim
                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-594572/56
                                                                                                                                                                                                                                                                                                                                                                                                                                               Berka RM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-MAR-2000; 2000WO-US07781.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Aspergillus niger.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAF11311 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (NOVO ) NOVO NORDISK BIOTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        selecting a composition for inhibiting prostate cancer in a patient; assessing the prostate cell carcinogenic potential of a compound; determining whether prostate cancer has metastasized in a patient;
                                                                                                                                                                                                                          87;
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                                                                                                                                                                                                                       Page 1716; 3161pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99US-0273623.
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                                                                                                                                                                                                                                                                               sequence
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                                                                                                                                                                                                                    English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      same genes in one or more second filamentous fungal cells. Monitoring the global expression of genes from FF cells allows the production potential of the microorganisms too be improved. New genes may be discovered, possible functions of unknown open reading frames can be identified and gene copy number variation and stability can be monitored. The expression of genes can be used to study how FF cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAH45451 standard;
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DB; AAG62454.
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/note= "Protein tyrosine phosphatase"
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This invention relates to a murine gene (Epm2A) and its encoded protein which cause epilepsy. Epm2A is a protein tyrosine phosphatase. Included in the invention is a method of preparing an Epm2A knockout mouse. The gene can be used for the analysis of Epm2A expression and to create an epilepsy model animal. The present sequence represents cDNA encoding

encoded protein hatase. Included

Claim

2; Page 6-8; l0pp; Japanese.

Mouse epilepsy-causing gene for the analysis mouse model of epilepsy -

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Best Local S
Matches 15
                                                         weight markers, as a food supplement, for generating antibodies, in medical imaging, screening and diagnostic assays and for treating cell-proliferative disorders (cancer), neurodegenerative diseases (Parkinson's or Alzheimer's disease), autoimmune diseases (multiple sclerosis, diabetes, hupus) genetic disorders, myeloid or lymphoid disorders, platelet or coagulation disorders, wound, burns, incision
                                                                                                                                                                                                                                                                                                                                                                                                                                   Tang YT,
Xue AJ,
            ulcers, liver or lung fibrosis, infections parasitic), arthritis, etc.
Note: The sequence data for this patent did
                                                                                                                                                   nucleotide sequence selected from any of 948 sequences (ABZ11119-ABZ12066) or their mature mature processing portion, active domain coding protein or complementary sequences. The polynucleotides are useful for identifying expressed genes or for physical mapping of human genome. The encoded polypeptides (ABP68902-ABP69849) are useful as molecular
                                                                                                                                                                                                                                                                                           New polynucleotides comprising sequences assembled from expressed sequence tags (ESTs), useful for treating cell-proliferative, neurodegenerative, autoimmune, genetic, myeloid or lymphoid, or platelet or coagulation disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; genome mapping; gene therapy; food supplement; virus; fungus; cell-proliferative disorder; neurodegenerative disease; bacterial; Parkinson's disease; Alzheimer's disease; autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 1419 BP;
                                                                                                                                                                                                                 The invention relates to an isolated polynucleotide (I) comprising nucleotide sequences
                                                                                                                                                                                                                                                              Claim 1; SEQ ID NO 707; 1012pp + Sequence Listing; English
                                                                                                                                                                                                                                                                                                                                                                                         WPI;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              arthritis; cytostatic; immunomodulator; nootropic; neuroprotective; antigarkinsonian; antidiabetic; immunosuppressive; dermatological;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              multiple sclerosis; diabetes; genetic disorder; wound; burn; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human polynucleotide SEQ ID
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T, Wang J,
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ence data for this patent did not form but was obtained in electronic format
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Ma Y, Yamazaki V, Chen R,
J, Wang D, Drmanac RT;
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R, Wang Z,
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Matches 15
                          The present sequence encodes C16N which can: (a) induce differentiati into cells which can degrade hydroxyapatite; (b) maintain neuron survival; (c) inhibit osteoblast proliferation; and (d) promote type collagen expression in osteoblasts. C16 and C16N are agents for the treatment of a broad range of diseases including hypercalcaemia, hypertension, diabetes, arteriosclerosis, myocardial infarction and terminal cancer. They may also be used as a screen for potential inhibitors of their activity for possible medicinal use. Transgenic animals containing DNA coding for the proteins can be used as model organisms and for the production of recombinant C16/C16N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  at
                                                                                                                                                                                                  Proteins C16 and C16N promote neuron survival and type 1 collagen production - for treatment of diseases involving collagen production, calcium regulation or neuron survival
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C16; C16N; neuron survival; type 1 collagen; calcium regulation; hypercalcaemia; hypertension; diabetes; arteriosclerosis; cancer
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                                                                                                                                                                           Claim 6; Page 61-62; 86pp; Japanese.
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P-PSDB; AAW31366.
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23-APR-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first
                                                                                                                                                                                                                                                                                               Mochizuki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    promoting neuron survival and type 1 collagen production.
                                                                                                                                                                                                                                                                                                                                                  97JP-0041562.
96JP-0127954.
                                                                                                                                                                                                                                                                                                                                                                                          97WO-JP01391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /*tag= a
/product= "C16N"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1..1740
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        512 A; 296 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CDNA; 1808
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   83.3%; Score 15;
100.0%; Pred. No.
tive 0; Mismatc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        287 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        604 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 24;
21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 1699;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           other;
                                                                                                                                                induce differentiation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
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Query Match
Best Local Similarity
Matches 15; Conserv

Conservative

0

Mismatches

0:

0;

Gaps

0

83.3%; Score 15; 100.0%; Pred. No.

DB 21; 18;

Length 1808;

1808

B₽;

336 A; 548

C; 551 G;

373

T; 0 other;

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RESULT 26
AAZ40179
ID AAZ40
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AAZ44730
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                                                                                                              Matches
                                                                                                                                      Query Match
                                                                                                                                                                                    promoter, or cartilage disorder treatment agent, which contains C16N, C16N-2; or their protein analogs, or one of the genes encoding them, as active ingredient. The promoters and cartilage disorder treatment agents are for the therapy of cartilage failure including deformans arthritis, diseases due to cartilage formation abnormality, deletion of cartilage in bone fraction, joint cartilage caused by injury, or damage to articular disc, acute purulent arthritis, tuberculous arthritis, syphilous arthritis, chronic rheumatoid arthritis, rheumatic fever, systemic erythematosus, deformative spinal diseases, and intervertebral hernia. This sequence encodes the human C16N-1 protein which is described in the method of the invention.
AAZ40179 standard; cDNA; 2293 BP.
                                                                                                                                                                 Sequence 2293 BP;
                                                                                                                                                                                                                                                                                                                                  This invention describes a novel cartilage cell differentiation
                                                                                                                                                                                                                                                                                                                                                          Example 2; Page 36-41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200001405-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cartilage disorder treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C16N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human C16N-1 cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAZ44730;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ44730 standard; cDNA; 2293
                                                                                                                                                                                                                                                                                                                                                                                               Cartilage cell differentiation promoters or remedies for cartilage
                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-126903/11.
P-PSDB; AAY51326.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ishiduka Y, Mochizuki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-JAN-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (SUMU ) SUMITOMO PHARM CO LTD
                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     949
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C16N-1; C16N-2; human;
                                                                                                             15;
                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GGCTGGTGTCACCTG
                                                                                        GGCTGGTGTCACCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GGCTGGTGTCACCTG
                                                               GGCTGGTGTCACCTG 1519
                                                                                                             Conservatjve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98JP-0190889.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        99WO-JP03577.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /*tag= a
/product= "C16N-1"
/note= "no stop codon given"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
107..2293
                                                                                                                                                               417 A; 716 C; 691 G; 469 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   entry)
                                                                                                                          83.3%;
                                                                                                                                                                                                                                                                                                                                                        55pp; Japanese
                                                                                       15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  963
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15
                                                                                                              0,
                                                                                                                            Score 15;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cartilage cell differentiation; therapy;
agent; arthritis; rheumatoid arthritis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP
                                                                                                                Mismatches
                                                                                                                             No.
                                                                                                                                        DB 21;
                                                                                                              0,
                                                                                                                                      Length 2293;
                                                                                                                Indels
                                                                                                              <u>;</u>
                                                                                                              Gaps
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RESULT 27
AAZ44731
ID AAZ44
XX
AC AAZ44
XX
DT 17-AF
DX Humar
XX
C16N;
KW C16N;
KW Carti
XX
CS Homo
XX
FT CDS
FT
YT CDS
FT
XX
W0201
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                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C16N-1; C16N-2; human; myeloid cell differentiation; neuron; hydroxyapatite decomposition; osteoblast growth inhibitor; type I collagen expression; ds.
                                                                                                                        C16N; C16N-1; C16N-2; human; cartilage cell differentiation; therapy; cartilage disorder treatment agent; arthritis; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 2293 BP; 417 A; 716 C; 691 G; 469 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            inducing differentiation of a mysloid cell to a cell having hydroxyapatite decomposing activity; (2) it has an activity of maintaining the survival of neuron; (3) it has an activity of inhibiting growth of osteoblasts; and (4) it has an activity of promoting expression of type I collagen in osteoblasts. The activity of the proteins can be limited to a specific tissue or a specific site.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human C16N-1 coding sequence
                                                                                                                                                                                             17-APR-2000
                                                                                                                                                                                                                         AAZ44731;
                                                                                                                                                                                                                                                 AAZ44731 standard; cDNA; 2301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This sequence encodes a C16N-1 protein of the invention. The invent also relates to C16N-2 proteins. The proteins of the invention have following features: (1) it has a differentiation inducing activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 2; Page 18-21; 30pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New proteins {\tt C16N-1} and {\tt C16N-2} or a gene coding them - can have activity limited to specific tissue or site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               JP11308995-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-FEB-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAZ40179;
                                                                                                                                                                  Human C16N-2 cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           P-PSDB; AAY55017
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-APR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (SUMU ) SUMITOMO SEIYAKU KK
                                                                                                                                                                                                                                                                                                                        1505 GGCTGGTGTCACCTG 1519
                                                                                              sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2000-046933/04.
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                                                                                                                                                                                                                                                                                                                                          GGCTGGTGTCACCTG 15
                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              98JP-0134440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98JP-0134440
                                                     Location/Qualifiers
95..2233
                          /product= "C16N-2"
                                                                                                                                                                                                                                                                                                                                                                                          83.3%; Score 15; 100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                              <u>,</u>
                                                                                                                                                                                                                                                    股
                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                          DB 21;
                                                                                                                                                                                                                                                                                                                                                                                                        21;
                                                                                                                                                                                                                                                                                                                                                                              0,
                                                                                                                                                                                                                                                                                                                                                                                                       Length 2293;
                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            promoting expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention ention have the
                                                                                                                                                                                                                                                                                                                                                                              0;
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WO200001405-A1.

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RESULT 28
AAZ40180
ID AAZ40
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     ACC XXX ACC XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C16N-1; C16N-2; human; myeloid cell differentiation; hydroxyapatite decomposition; osteoblast growth inhit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAZ40180 standard; cDNA; 2301 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 2301 BP; 425 A; 714 C; 703 G; 459 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     treatment agents are for the therapy of cartilage failure including deformans arthritis, diseases due to cartilage formation abnormality, deletion of cartilage in bone fraction, joint cartilage caused by injury, or damage to articular disc, acute purulent arthritis, tuberculous arthritis, chronic rheumatoid arthritis, rheumatic fever, systemic erythematosus, deformative spinal diseases, and intervertebral hernia. This sequence encodes the human C16N-2 protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention describes a novel cartilage cell differentiation promoter, or cartilage disorder treatment agent, which contains C16N, C16N-1, C16N-2, or their protein analogs, or one of the genes encodir them, as active ingredient. The promoters and cartilage disorder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2000-126903/11.
P-PSDB; AAY51327.
New proteins C16N-1 and C16N-2 or a gene coding them -
                                                     P-PSDB;
                                                                                                                                                                                                                                                                                                                                                       JP11308995-A.
                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human C16N-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAZ40180;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cartilage cell differentiation promoters or remedies for cartilage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-JAN-2000
                                                                                                                                                                                            28-APR-1998;
                                                                                                                                                                                                                                               28-APR-1998;
                                                                                                                                                                                                                                                                                                    09-NOV-1999.
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                                                                                                                                       (SUMU ) SUMITOMO SEIYAKU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1442 GGCTGGTGTCACCTG 1456
                                                     2000-046933/04.
DB; AAY55018.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               collagen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 GGCTGGTGTCACCTG 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Page 45-50;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mochizuki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       coding sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 98JP-0190889.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               expression; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          83.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   55pp; Japanese
                                                                                                                                       즛
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 15; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 21;
21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           growth inhibitor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    neuron;
  can have
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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RESULT 29
AAZ33584
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local
                         This invention describes novel human nucleic acid sequences from normal breast tissue which have cytostatic activity. The nucleic acid sequences can be used to produce and isolate full-length gene sequences. They can be used to express proteins, which can be used as tools to find an activity against breast cancer. The sequences can be used in sense or antisense form. They are especially useful for medicaments for gene therapy to treat breast cancer and for treating illnesses associated with fat metabolism. AAZ33541-Z33610 represent expressed sequence tags described in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         inducing differentiation of a myeloid cell to a cell having hydroxyapatite decomposing activity; (2) it has an activity of maintaining the survival of neuron; (3) it has an activity of inhibiting growth of osteoblasts; and (4) it has an activity of promoting expression of type I collagen in osteoblasts. The activity of the proteins can be limited to a specific tissue or a specific site.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 2301 BP; 425 A; 714 C; 703 G; 459 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This sequence encodes a C16N-2 protein of the invention. The invention also relates to C16N-1 proteins. The proteins of the invention have the following features: (1) it has a differentiation inducing activity,
                                                                                                                                                                                 Claim 3; 132-133; 206pp; German.
                                                                                                                                                                                                              Human nucleic acid sequences and protein products from normal breast tissue, useful for breast cancer therapy
                                                                                                                                                                                                                                                            WPI; 1999-528979/45.
                                                                                                                                                                                                                                                                                        Specht T,
                                                                                                                                                                                                                                                                                                                                                   20-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                 20-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                               23-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                              DE19813835-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        medicaments;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Expressed sequence tag; EST; human; breast; cancer; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human breast tumour-associated EST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-DEC-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ33584;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAZ33584 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 2; Page 23-26; 30pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              activity limited to
                                                                                                                                                                                                                                                                                                                        (META-) METAGEN GES GENOMFORSCHUNG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1442 GGCTGGTGTCACCTG 1456
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 GGCTGGTGTCACCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                        Hinzmann B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gene therapy; treatment; fat metabolism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first
                                                                                                                                                                                                                                                                                                                                                     98DE-1013835.
                                                                                                                                                                                                                                                                                                                                                                                   98DE-1013835
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              specific tissue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 83.3%;
                                                                                                                                                                                                                                                                                        Schmitt A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2952
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 15;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              얁
                                                                                                                                                                                                                                                                                        Pilarsky C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB
21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21;
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                                                                                                                                                                                                                                                                                          Dahl
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          88
                                                                                                                                                                                                                                                                                            Rosenthal A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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Query Match

Sequence 2952 BP; 925 A; 564

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881 T; 0 other;

Best Local Similarity

83.3%;

Score 15; Pred. No. 582 G;

DB 20; 20;

Length 2952;

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ARESULT 30
AAC777
XX AAC77
CCC AAC7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antidiabetic; antiasthmatic; antirheumatic; antiarthritic; antiviral; antiinflammatory; antithyroid; antiallergic; antibacterial; cardiant; dermatological; neuroprotective; thrombolytic; coagulant; nootropic; vasotropic; antipsoriatic; antiangiogenic; gene therapy; inflammation; immune disorder; haematopoietic cell disorder; autoimmune disorder; allergic reaction; graft versus host disease; organ rejection; haemostatic; thrombolytic; cardiovascular disorder; infection;
                                                                                         Polynucleotides, polypeptides, antibodies, agonists and antagonists from the present invention may be used to treat immune disorders by activating or inhibiting the proliferation, differentiation or mobilisation of immune cells, to treat disorders of haematopoietic cells, autoimmune disorders, allergic reactions, graft versus host disease and organ rejection, modulate haemostatic or thrombolytic activity, modulate inflammation, cancers, cardiovascular disorders, neurological disease and bacterial or viral infections. The peptides, mucleotides, antibodies, agonists and antagonists may be also be used in drug screens. AAC78449 to AAC78457 and AAB44240 represent sequences used in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                            antidiabetic; antiasthmatic; antirheumatic; antiarthritic; antiinflammatory; antithyroid; antiallergic; antibacterial; antiviral; dermatological; neuroprotective; cardiant; thrombolytic; coagulant; nootropic; vasotropic; antipsoriatic and antiangiogenic. The polynucleotides and polypeptides can be used for preventing, treating o ameliorating medical conditions and diagnosing pathological conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel isolated nucleic acids comprising sequences encoding peptides useful for treating or diagnosing {\rm e.g.} cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAC77831 standard; cDNA; 3035
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 796-797; 2352pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200055350-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        neurological disease; drug screening; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human cancer associated gene sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAC77831;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rosen CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-MAR-2000; 2000WO-US05882.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            tissues and cells the genes are expressed in. Example of activities include: cytostatic; proliferative; vulnerary; immunomodulator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TO AAB43398 to AAB44239. The proteins can have activities based on the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAC77607 to AAC78448 encode the human cancer associated proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2836
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2000-587533/55
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ruben SM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cytostatic; proliferative; vulnerary;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99US-0124270
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO:225.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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Sequence 3035 BP; 955 A;

575 C;

588

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912 .. ហ

other,

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CTGGTGTCACCTGTT 17

Matches Query Match Best Local

15;

Conservative

0;

Similarity

100.0%; 83.3%;

Score 15; Pred. No. Mismatches

20;.

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Indels

0

Gaps

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DB 22;

Length 3035;

Sequence 3035 BP; 955 A; 575 C; 590 G;

912 T; 3 other;

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RESULT 31
AAH34809
ID AAH34
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Best Local Similarity
Matches 15; Conserv
                                                                                      cancer antigens have cytostatic activity and can be used in gene therapy and vaccine production. N and P may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate P expression. For example, N and P may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of P by expressing inactive proteins or to supplement the patients own production of P. Additionally, N may be used to produce the colon cancer-associated Ps, by inserting the nucleic acids into a host cell and culturing the cell to express the proteins. N and P can be used in the prevention, diagnosis and treatment of colorectal carcinomas and cancers. AAH37196 to AAH37204 and AAB77789 represent sequences used in the exemplification of the
                          N.B. Pages 666 to 682 and page 7053 of the sequence listing were missing at time of publication, meaning no sequences are present SEQ ID NO:1027 to 1052, 7921 and 7922.
                                                                                                                                                                                                                                                                                                                                                                                      Nucleic acids encoding 4277 human colon cancer-associated polypeptides, useful for preventing, diagnosing and/or treating colorectal cancers -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAH34809 standard; cDNA; 3035
                                                                             present invention.
                                                                                                                                                                                                                                                                                         the proteins are collectively known as colon cancer antigens. The colon
                                                                                                                                                                                                                                                                                                        cancer-associated nucleic
                                                                                                                                                                                                                                                                                                                         AAH32943 to AAH37195 and
                                                                                                                                                                                                                                                                                                                                                      Claim 1; Page 3405-3406; 9803pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-235357/24.
P-PSDB; AAG75404.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ruben SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-SEP-1999;
03-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-SEP-2000; 2000WO-US26524
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-APR-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      colorectal carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; colon cancer; colon cancer antigen; diagnosis; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human colon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     03-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAH34809
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2771 CTGGTGTCACCTGTT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Barash SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cancer antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        99US-0157137.
99US-0163280.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              83.3%; Score 15; 100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Birse CE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2785
                                                                                                                                                                                                                                                                                                       AAG73514 to AAG77788 represent human colon c acid molecules (N) and proteins (P), where
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   encoding cDNA SEQ ID NO:1891.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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                                             present
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                                                                           RESULT 33
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                                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence encodes C16N which can: (a) induce differentiation into cells which can degrade hydroxyapatite; (b) maintain neuron survival; (c) inhibit osteoblasts proliferation; and (d) promote type 1 collagen expression in osteoblasts. C16 and C16N are agents for the treatment of a broad range of diseases including hypercalcaemia, hypertension, diabetes, arteriosclerosis, myocardial infarction and terminal cancer. They may also be used as acreen for potential inhibitors of their activity for possible medicinal use. Transgenic animals containing DNA coding for the proteins can be used as model
          AAZ44729 standard; cDNA; 3337 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Proteins C16 and C16N promote neuron survival and type 1 collagen production - for treatment of diseases involving collagen production, calcium regulation or neuron survival
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C16; C16N; neuron survival; type 1 collagen; calcium regulation; hypercalcaemia; hypertension; diabetes; arteriosclerosis; cancer; myocardial infarction; hydroxyapatite; osteoblast; ds.
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                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                   organisms and for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ishiduka Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-OCT-1997.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C16N gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAV02312;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (SUMU ) SUMITOMO PHARM CO LTD
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                                                                                                                                                 1184 GGCTGGTGTCACCTG 1198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              6; Page 55-57; 86pp; Japanese.
                                                                                                                                                                                                  1 GGCTGGTGTCACCTG 15
                                                                                                                                                                                                                                                    l Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAW31364.
                                                                                                                                                                                                                                                                                                                                                 3065 BP; 571 A; 950 C; 889
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                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mochizuki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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96JP-0127954.
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/product= "C16N"
                                                                                                                                                                                                                                                                                                                                                                                                   the production of recombinant
                                                                                                                                                                                                                                                 83.3%; Sc
100.0%; F
ative 0;
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                                                                                                                                                                                                                                                                           Score 15;
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                                                                                                                                                                                                                                                                                                                                                                                                            This invention describes a novel cartilage cell differentiation promoter, or cartilage disorder treatment agent, which contains C16N, C16N-1, C16N-2, or their protein analogs, or one of the genes encoding them, as active ingredient. The promoters and cartilage disorder treatment agents are for the therapy of cartilage failure including deformans arthritis, diseases due to cartilage formation abnormality, deletion of cartilage in bone fraction, joint cartilage caused by injury, or damage to articular disc, acute purulent arthritis, tuberculous arthritis, syphilous arthritis, chronic rheumatoid arthritis, rheumatic fever, systemic erythematosus, deformative spinal diseases, and intervertebral hernia. This sequence encodes the murine C16N-2 protein that the cartical intervertebral hernia. This sequence encodes the murine C16N-2 protein
C16N-1; C16N-2; mouse; myeloid cell differentiation; neuro, hydroxyapatite decomposition; osteoblast growth inhibitor; type I collagen expression; ds.
                                                              Mouse C16N-2 coding sequence
                                                                                                 22-FEB-2000
                                                                                                                              AAZ40178;
                                                                                                                                                            AAZ40178 standard;
                                                                                                                                                                                                                                                                                                                                                                   Sequence 3337 BP; 633 A; 1037 C; 976 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Page 27-33; 55pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cartilage cell differentiation promoters or remedies for cartilage
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C16N; C16N-1; C16N-2; murine; cartilage cell differentiation; therapy; cartilage disorder treatment agent; arthritis; rheumatoid arthritis; s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Murine C16N-2 cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-APR-2000
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                                                                                                                                                                                                                                                                                                                                                                                                  is described in the method of the
                                                                                                                                                                                                                                                                        1 GGCTGGTGTCACCTG
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                                                                                                                                                                                                                                           GGCTGGTGTCACCTG
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/product= "C16N-2"
                                                                                                                                                             CDNA; 3337
                                                                                                 entry)
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                                                                                                                                                                                                                                                                                                                        83.3%; Score 15;
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Matches 15
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                                                                                                                                                    02-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-APR-2000 (first entry)
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Ishiduka Y, Mochizuki R;
                                                                                                          06-JUL-1998;
                                                                                                                                                                                                                         13-JAN-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C16N; C16N-1; C16N-2; murine; cartilage cell differentiation; cartilage disorder treatment agent; arthritis; rheumatoid arth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Murine C16N-1 cDNA.
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                                                      (SUMU ) SUMITOMO PHARM CO LTD
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                                                                                                            98JP-0190889.
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395..2584
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/product= "C16N-1"
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100.0%; Pr
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Pred. No.
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                                                                                                                                                     Sequence 3674 BP; 695 A; 1146 C; 1044 G; 789 T; 0 other;
                                                                                                                                                                                             which
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 2; Page 18-24; 55pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cartilage cell differentiation promoters or remedies for cartilage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2000-126903/11.
P-PSDB; AAY51324.
1793 GGCTGGTGTCACCTG
                                                                                                                                                                                           is described in the method of the invention.
                        GGCTGGTGTCACCTG
                                                                            Conservative
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1807
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/cgn2_6/ptodata/2/ina/6B_COMB.seq:*
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                                                                                                                                TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-433-4150
TELEPAX: 415-433-8716
INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 29 Application US/08197791
Patent No. 15556772 Application US/08197791
GENERAL INFORMATION:
APPLICANT: Worge, Joseph A.
APPLICANT: Wullinax, Rebecca L.
TITLE OF INVENTION: NOVEL POLYMERASE COMPOSITIONS AND USES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         110
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APPLICATION NUMBER: US 08/
FILING DATE: 08-DEC-1993
ATTORNEY/AGENT INFORMATION:
NAME: BOTTINEY, SCOTT R.
REGISTRATION NUMBER: 34,29
REFERENCE/DOCKET NUMBER: S
                                             MOLECULE TYPE: HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
CONFUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
                                                                                TYPE: STRANDEDNESS: SIN
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US-09-1
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US-08-632-470-50
US-07-936-163-2
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US-09-107-532A-2348
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US-08-964-494-3
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US-09-107-532A-
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-107-532A-2348
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Query Match

100.0%;

Score 20;

DB 1; Length 20;

CORRESPONDENCE ADDRESS:
ADDRESSEE: Cooley Godward Castro Huddleson & Tatum

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RESULT 3
PCT-US95-04611A-18
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US-08-917-320-18
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Sequence 18, Application PC/TUS9504611A GENERAL INFORMATION:
APPLICANT: Spacte, Richard and Jackman, TITLE OF INVENTION: Non Splicing Variant NUMBER OF SEQUENCES: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 18 Patent No.
                                                                                                                                                                                                                              Query Match
Best Local Similarity
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TELEX: 380816 CooleyPA
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 08/229,29
FILING DATE: April 18, 1994
ATTORNEY/AGENT INFORMATION:
NAME: LUAIN CSETT
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: AVIR
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtent In Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/917,320
FILING DATE: 25-AUG-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                             FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Spaete, Richard and Jackman, Winthrop, TITLE OF INVENTION: No. 5824508 Splicing Variants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                              NAME/KEY:
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STATE: California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: 415-843-5163
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                                                                                                                                                 2899 CCTTAGGAGGAACAAGTCCC 2918
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o. 5824508
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EDNESS: double
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Pred. No. 0.0089;
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                ackman, Winthrop, T.
Variants of gp350/220
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RESULT 4
US-08-783-774-1
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Best Local Similarity
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ZIP: 94306
ZIP: 94306
COMPUTER REALDABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: PC-DOS/MS-DOS
TYPEN: PC-DOS/MS-DOS
TYPEN: PC-DOS/MS-DOS
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TELEX: 380816 CooleyPA
INFORMATION FOR SEQ ID NO:
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            SOFTWARE: FastSEQ Version 2.0 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/783,774 FILING DATE: 15-JAN-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: LUBIN CGERT
REGISTRATION UNMER: 31,822
REFERENCE/DOCKET NUMBER: AVIR-003/00US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-843-5163
TELEFAX: 415-857-0663
                                                                                                                                                                                                            NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                              APPLICANT: Spaete, Richard APPLICANT: Jackman, Winthrought OF INVENTION: NON-SPITITLE OF INVENTION: GP350/:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
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                                                                        COMPUTER: IBM CON
OPERATING SYSTEM:
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LOCATION: 1014..3734
                                                                                                             MEDIUM TYPE:
                                                                                                                                               ZIP:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
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                                                                                                                                                          COUNTRY:
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CLASSIFICATION:
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R READABLE FORM:
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                                                                                                                                                               USA
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VENTION: NON-SPLICING VARIANTS OF
VENTION: GP350/220
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                                                                                             IBM Compatible
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Pred. No. 0.0089;
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US-09-620-312D-823

Patent No. 656966

Application US/09620312D

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GENERAL

APPLICANT: Tang, Y. Tom
APPLICANT: Liu, Chenghua

APPLICANT:

Asundi, Vin Zhang, Jie

RESULT 6

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Query Match
Best Local Similarity
Thehes 20; Conserva
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                                                                                                                                         ; OTHER INFORMATION: gp350/220 US-09-556-706B-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-556-706B-1
                                                                                                                                                                                                                             SOFTWARE: PatentIn version 3.0 SEQ ID NO 1 LENGTH: 5931
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APPLICANT: Spaete, Richard
APPLICANT: Jackman, Winth
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                                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 1994-04-18
NUMBER OF SEQ ID NOS: 19
                                                                                                                                                                                                                                                                                                                      PRIOR FILING DATE: 1997-01-15
PRIOR APPLICATION NUMBER: 08/229,291
                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: 7682-050-999
CURRENT APPLICATION NUMBER: US/09/556,706B
CURRENT FILING DATE: 2000-04-24
PRIOR APPLICATION NUMBER: 08/783,774
PRIOR APPLICATION NUMBER: 08/783,774
                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Jackman, Winthrop
TITLE OF INVENTION: NON SPLICING VARIANTS OF GP350/220
                                                                                                                                                                                        TYPE: DNA
ORGANISM: Virus
                                                                                                                                                                             FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: 212-790-9090
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2899 CCTTAGGAGGAACAAGTCCC 2918
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                212-869-8864
                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
                                                                                 100.0%; Score 20; DB 4; 100.0%; Pred. No. 0.0087;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
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                                                                    <u>,</u>
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Pred. No. 0.0087;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 3; Length 5931;
                                                                                                  DB 4; Length 5931;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
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                                                                    <u>.</u>
                                                                 Gaps
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Feiyan

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CURRENT APPLICATION NUMBER: US/09/620,312D
CURRENT FILING DATE: 2000-07-19
PRIOR APPLICATION NUMBER: 09/52,317
PRIOR FILING DATE: 2000-04-25
PRIOR APPLICATION NUMBER: 09/488,725
PRIOR FILING DATE: 2000-01-21
PRIOR FILING DATE: 2000-01-21
VUMBER OF SEQ ID NOS: 1105
SOPTWARE: pt_FL_genes Version 1.0
SEQ ID NO 823
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                                                                                                                      US-09-439-313-354
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US-09-439-313-354
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                                                                                                                                         TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER FILE REFERENCE: 210121.427C9
CURRENT APPLICATION NUMBER: US/09/439,313
CURRENT FILING DATE: 1999-11-12
NUMBER OF SEQ ID NOS: 575
SOFTWARE: FASESEQ for Windows Version 3.0
SEQ ID NO 354
LENGTH: 854
TYPE: DNA
ORGANISM: HOMO Sapien
Query Match
Best Local Similarity
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APPLICANT: Dil
APPLICANT: Mit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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APPLICANT: Wang, Dunrui
APPLICANT: Wang, Zhiwei
APPLICANT: John Tillinghast
APPLICANT: Jornanac, Radoje T
TITLE OF INVENTION: No. $569662el Nucleic Acids and
TITLE OF INVENTION: Polypeptides
FILE REFERENCE: 788CIPED
FILE RE
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LOCATION: (521)..(2611)
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TYPE: DNA
ORGANISM: Homo sapiens
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Yang, Yonghong
Wang, Jian-Rui
Zhou, Ping
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Solk, John
Day, Craig
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Reed, Steven G.
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Mitcham, Jennifer L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Retter, Mark
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Harlocker,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Jiangchun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  John
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   70.0%;
100.0%;
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       Score 14;
Pred. No.
       DB 4;
20;
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SEQ ID NO 354
LENGTH: 854
TYPE: DNA
ORGANISM: Homo sapien
US-09-352-616A-354
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           APPLICATION NUMBER: US/08/599,171A
FILING DATE: Concurrently
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
CLASSIFICATION: 435
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: HERRON, CHARLES J.
REGISTRATION NUMBER: 28,019
REFERENCE/DOCKET NUMBER: 331400-38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Dillon, Davin C.
APPLICANT: Harlocker, Susan Louise
APPLICANT: Jiang, Yuqui
APPLICANT: Xu, Jiangchun
APPLICANT: Mitcham, Jennifer Lynn
APPLICANT: Mitcham, Jennifer Lynn
TITLE OF INVENTION: COMPROSTATE CANCER AND METHODS FOR THEIR USE
TITLE OF INVENTION: OF PROSTATE CANCER AND METHODS FOR THEIR USE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: 210121.427C8
CURRENT APPLICATION NUMBER: US/09/352,616A
CURRENT FILING DATE: 1999-07-13
                                                                                                                                                                                                                           OPERATING SYSTEM: MS-DOS
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 INCH DISKETTE
COMPUTER: IBM PS/2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: WARREN, PATRICK V.
TITLE OF INVENTION: TRANSAMINASES AND AMINOTRANSFERASES
NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                               CITY: ROSELAND
STATE: NEW JERSEY
                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: CARELLA, BYRNE, BAIN, GIL
ADDRESSEE: CECCHI, STEWART & OLSTEIN
STREET: 6 BECKER FARM ROAD
                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
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o. 5814473
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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100.0%; Pred. No.
Live 0; Mismatc
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TELEPHONE:

201-994-1700

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RESULT 10
US-08-646-590B-17/c
                                                  US-08-646-590B-17
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Query Match
Best Local Similarity
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                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 201-994-1744
NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/5:
FILING DATE: 09-FEB-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/646,590B
FILING DATE: 08-May-1996
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Warren, P
APPLICANT: Swanson,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: GENOMIC DNA
                                                                                                          MOLECULE TYPE:
                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION: TELEPHONE: 619/678-5070
                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Halle, Ph.D., Lisa A.
REGISTRATION NUMBER: 38,347
REFERENCE/DOCKET NUMBER: 09
                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US97/01094
FILING DATE: 21-January-1997
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Match 70.0%;
Local Similarity 100.0%;
                                                                              NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM CONTROL OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: UZIP: 92037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 4225 Executive Square, Suite 1400 CITY: La Jolla
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: NUCLEIC ACID
                                                                                                                                        STRANDEDNESS:
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                                                                                                                             ropology:
                                                                 OCATION:
                                                                                                                                                                          ENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14; Conservative
                                                                                                                                                         H: 1245 base pairs nucleic acid
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Swanson, Ronald V.
                                                             Coding Sequence 1...1242
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                                                                                                                              linear
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                                                                                                              Genomic DNA
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                                                                                                                                          single
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70.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TRANSAMINASES AND AMINOTRANSFERASES
                                                                                                                                                                                                                                                                                                                                                                                                                     08/599,171
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 Score 14;
; Pred. No.
                                                                                                                                                                                                                                                                         09010/017001
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Pred. No
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 DB 2;
19;
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                Length 1245;
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                                                                                                                                             RESULT 12
US-09-412-184-17/c
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                                                                                                  Sequence 17, Application US/09412184
Patent No. 6268188
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                   Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17, Application US/09069226
Patent No. 6013509
                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 201-994-1744
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: HERRON, CHARLES J.
REGISTRATION NUMBER: 28,019
REFERENCE/DOCKET NUMBER: 33:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: PRIOR APPLICATION NUMBER:
                                           APPLICANT: Warren, Patrick V. APPLICANT: Swanson, Ronald V. TITLE OF INVENTION: TRANSAMINASES NUMBER OF SEQUENCES: 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: GENOMIC DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: CARELLA, BYRNE, BAIN, GIL
ADDRESSEE: CECCHI, STEWART & OLSTEIN
STREET: 6 BECKER FARM ROAD
                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS: SI
TOPOLOGY: LINEAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: ROSELAND
STATE: NEW JERSEY
   STREET:
                                                                                                                                                                                                                                                                                                                                                                                         TYPE: NUCLEIC ACID
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               ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                          ENGTH:
                                                                                                                                                                                                            1069 TTAGGAGGAACAAG 1056
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                                                                                                                                                                                                                                        3 TTAGGAGGAACAAG 16
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E: Fish & Richardson, P.C.
4225 Executive Square, Sui
                                                                                                                                                                                                                                                                                                                                                                                                        1245 NUCLEOTIDES
                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                                  70.0%; Score 14; 100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08/599,171
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 Suite 1400
                                                           AND AMINOTRANSFERASES
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COUNTRY: U

CA

Query Match

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US-09-232-160-6/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
OTHER INFORMATION: 1698542 - 09-232-160-6
                                                                                                                                                                                                      TITLE OF INVENTION: DETECTION OF ALTERED EXPRESSION OF GENES REGULATING CELL TITLE OF INVENTION: PROLIFERATION FILE REFERENCE: PA-0003 US
                                                                                                                                                                                                                                                          APPLICANT: James Gilmore
APPLICANT: Susan G. Stuart
APPLICANT: Laura Stuve
                                                                                                                            NUMBER OF SEQ ID NOS: 23
SOFTWARE: PERL Program
                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/232,160
CURRENT FILING DATE: 1999-01-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: Haile, Ph.D., Lisa A.
REGISTRATION UNMBER: 38,347
REFERENCE/DOCKET NUMBER: 0901
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619/678-5070
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INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                    APPLICANT: Steve Daniel
                                  TYPE: DNA
ORGANISM: Homo sapiens
FEATURE: -
                                                                                         ENGTH: 1585
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 09-FEB-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US97/01094
FILING DATE: 21-January-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/646,590
FILING DATE: 08-May-1996
APPLICATION NUMBER: 08/599,171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
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TOPOLOGY: lir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 1245 base pairs
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US-08-480-784-21/c
; Sequence 21, Application US/08480784
; Patent No. 5693473
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                                                                                                                                                                                                                             PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/300,266

FILING DATE: 02-SEP-1994

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/289,221

FILING DATE: 12-AUG-1994

ATTORNEY/AGENT INFORMATION:
                                                                                                                TELEFAX: 202-962-8300 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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FILING DATE: 29-NOV-1994
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 08/308,104
                                                                                                SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 24-MAR-PRIOR APPLICATION DATA:
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PRIOR APPLICATION DATA:
US 08/409,305
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                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
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HYPOTHETICAL:
               MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: FILING DATE: CLASSIFICATION: 435
                                                                                                                                                                                                 NAME: Ihnen, Jeffrey L. REGISTRATION NUMBER: 28,957
                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 0: FILING DATE: 24-MAR-1995
                               TOPOLOGY:
                                             STRANDEDNESS:
                                                                                                                                                 TELEPHONE:
                                                                                                                                                                                   REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 16-SEP-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Washington
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                                                                nucleic acid
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                                                                           4249 base pairs
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Futreal,
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                          linear
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Tavtigian, Sean V.
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               DNA (genomic)
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17q-Linked Breast and Ovarian Cancer
Susceptibility Gene
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; ORIGINAL SOURCE:
; ORGANISM: Hom
US-08-480-784-21
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Best Local Similarity
TELEFAX: 202-962-8300
NFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF
                             NAME: Ihnen, Jeffræy L.
REGISTRATION NUMBER: 28,957
REFERENCE/DOCKET NUMBER: 24
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-962-4810
                                                                                                                                                      FILING DATE: 02-SEP-
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                FILING DATE: 29-NOV-PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/409,305
                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                         FILING DATE: 12-AUG-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                   PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                  APPLICATION NUMBER: US 08/308,104 FILING DATE: 16-SEP-1994
                                                                                                                                                                                                                                                                                                                 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: U
ZIP: 20005
                                                                                                                                                                                      APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 435
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                                                                                                                                       APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             E: Venable, Baetjer, Howard & Civiletti, LLP
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Tavtigian, Sean V.
Wiseman, Roger W.
Futreal, P. Andrew
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Goldgar, David E.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                 24-MAR-1995
                                                                                                                                                                    02-SEP-1994
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100.0%; Pr
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                                                                                                                                       US 08/289,221
                                                                                                                                                                                      US 08/300,266
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; Pred. No. 18;
                                                             24884-109347
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RESULT 16
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Patent No.
                                                         FILING DATE: 02-SEP-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/289
FILING DATE: 12-AUG-1994
ATTORNEY/AGENT INFORMATION:
NAME: Ihnen, Jeffrey L.
REGISTRATION NUMBER: 28,957
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ANTI-SENSE: NO
ORIGINAL SOURCE:
                                                                                                                                                                                          FILING DATE: 16-SEP-
PRIOR APPLICATION DATA:
                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 24
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-962-4810
                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/409,305
FILING DATE: 24-MAR-1995
                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/487,002 FILING DATE:
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                                                                                                                                                                             APPLICATION NUMBER: US 08/300,266
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Homo sapiens
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202-962-8300
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Durocher, Francine
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                                                                                                                              US 08/289,221
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                                                24884-109347
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; ANTI-SENSE: NO; ORIGINAL SOURCE: ; ORIGINAL SOURCE: ; ORGANISM: Homo Bapiens US-08-487-002-21
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Best Local Similarity
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                                                                                                                                                                                                                 CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/
PILING DATE: 24-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/
FILING DATE: 29-NOV-1994
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HYPOTHETICAL: NO
HYPOTHETICAL: NO
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APPLICANT:
APPLICANT:
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                                                                        APPLICATION NUMBER: US 08/300,266
FILING DATE: 02-SEP-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/289,221
FILING DATE: 12-AUG-1994
                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                       FILING DATE: 12-AUG-1994
ATTORNEY/AGENT INFORMATION:
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 TELECOMMUNICATION INFORMATION:
                                                                                                                                                      PRIOR APPLICATION DATA:
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APPLICATION NUMBER: US 08/308,104
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
REFERENCE/DOCKET NUMBER: 248
                             NAME: Ihnen, Jeffrey REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/01 FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid-
STRANDEDNESS: double
TOPOLOGY: linear
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1201 New York Avenue, N.W., Suite 1000
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Tavtigian, Sean V.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Wiseman, Roger W
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              24884-109347
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Pred. No.
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18;
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RESULT 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Patent No.
                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/409,305
FILING DATE: 24-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/348,824
FILING DATE: 29-NOV-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/308,104
FILING DATE: 16-SEP-1994
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                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/488,011B
FILING DATE: 07-JUN-1995
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            FILING DATE: 02-SEP-1994
PRIOR APPLICATION DATA:
                                                             FILING DATE: 16-SEP-PRIOR APPLICATION DATA:
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MEDIUM TYPE: Floppy
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                                           APPLICATION NUMBER:
                                                                                                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Homo sapiens
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1201 New York Avenue, N.W., Suite 1000
                                                                                                                                                                                                                                                                                                                                                                                                         USA
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                                                                                                                                                                                                                                                                                                                                                      Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17q-Linked Breast and Ovarian Cancer
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                                                US 08/300,266
US 08/289,22:
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18;
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                                             FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/483,554
FILING DATE: 07-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/348,824
FILING DATE: 29-NOV-1994
PRIOR APPLICATION DATA:
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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MOLECULE TYPE: IC
HYPOTHETICAL: NO
ANTI-SENSE: NO
ORIGINAL SOURCE:
ORGANISM: Home
                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSE: Venable, Baetjer, Howard & Civiletti, LLP
ADDRESSE: Venable, Baetjer, Howard & Civiletti, LLP
                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IMP PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/850,727
                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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               APPLICATION NUMBER: US 08/308,104 FILING DATE: 16-SEP-1994
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APPLICATION DATA:
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Shattuck-Eidens, Donna M.
Tavtigian, Sean V.
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18;
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                                                                                                                                                                                          ZIP: 20005
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/10202
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LENGTH: 4249 base pair
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ATTORNEY/AGENT INFORMATION:
                                           APPLICATION NUMBER: US 0 FILING DATE: 24-MAR-1995 PRIOR APPLICATION DATA:
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REFERENCE/DOCKET NUMBER: 24/
TELECOMMUNICATION INFORMATION:
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APPLICATION NUMBER: US 08/289,221
FILING DATE: 12-AUG-1994
                                                                                            PRIOR APPLICATION DATA:
                                                                                                           PRIOR APPLICATION DATA:
APPLICATION.NUMBER: US
FILING DATE: 07-JUN-1995
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                              STREET: 12v.
CITY: Washington
PRIOR APPLICATION DATA:
                                                                                                                                                            FILING DATE:
CLASSIFICATION:
            APPLICATION NUMBER: US 08/348,824 FILING DATE: 29-NOV-1994
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1201 New York Avenue, N.W., Suite 1000
                                                                                                                                                                                                                                                                                                                                USA
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Durocher, Francine
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                                                                              US 08/409,305
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Pred. No.
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APPLICATION NUMBER:

16-SEP-1994

US 08-308,104

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PILING DATA:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US
APPLICATION O7-JUB-1995
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APPLICATION NUMBER: US 08/
FILING DATE: 12-AUG-1994
ATTORNEY/AGENT INFORMATION:
NAME: Ihnen, Jeffrey L.
REGISTRATION NUMBER: 28,95
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                                                                                         SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/10203
                                                                                                                                       CONPUTER READABLE FORM:
NEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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TYPE: nucleic acid
STRANDEDNESS: double
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Shattuck-Eidens, Do
Tavtigian, Sean V.
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Kamb, Alexander
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Goldgar, David E.
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GENERAL INFORMATION:
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Best Local
COUNTY...
ZIP: 2005
ZIP: 2005
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Flop
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NFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: DI
HYPOTHETICAL: NO
ANTI-SENSE: NO
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08
FILING DATE: 02-SEP-1994
PRIOR APPLICATION DATA:
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FILING DATE: 12-AUG-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
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                                                                                                                                                                                                                                                                      CITY: Washington
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REGISTRATION NUMBER: 28
                                                                                                                                                                                                                                                                                                                                             ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE:
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Shattuck-Eidens, D.
Tavtigian, Sean V.
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Goldgar, David E.
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          PatentIn Release #1.0, Version #1.30
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P. Andrew
Method for Diagnosing a
Method for Diagnosing a
Method for Diagnosing a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Roger W
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; Pred. No. 18;
0; Mismatches
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CURRENT APPLICATION DATA:

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US-08-746-411A-9/c
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Best Local
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                                                                                                                                                                                                                                                      GENERAL INFORMATION:
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SEQUENCE CHARACTERISTICS:
       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
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HYPOTHETICAL: NO
ANTI-SENSE: NO
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ORGANISM: Hom
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                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                           APPLICANT: O'MAHONY, TITLE OF INVENTION: 1
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APPLICATION NUMBER:
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CURRENT APPLICATION DATA:
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                                                                                                   COUNTRY:
                                                                                                                                 CITY: Gainesville
                                                                                                                                               STREET:
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1300 Gould Drive
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                                                                                                                                                             Ph.D., Esq.
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               Version #1.30
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18;
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                                                                                                                                                             SOFTWARE: Patentin Release #1.0, V.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/857,046A
FILING DATE: 15-MAY-1997
CLASSIFICATION: 435
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PRIOR APPLICATION DATA:
US 60/006461
                                APPLICATION NUMBER: 10-NOV-1995
FILING DATE: 10-NOV-1995
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 08/746,411
                                                                                                                                                                                                                                                                           ZIP: 30504
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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APPLICATION NUMBER: US 60/006461
FILING DATE: 10-NOV-1995
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            FILING DATE: 08-NOV-1996 PRIOR APPLICATION DATA:
                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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OPERATING SYSTEM: PC-DOS/MS-DOS
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APPLICATION NUMBER:
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1300 Gould Drive
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Alverez, Vernon L
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/desc = "individual isolate"
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100.0%; Pr
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Tissues and Methods of Identifying and Using the Same
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WO PCT/IE96/00073
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Pred. No.
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                                                                                                                                                                                                                                 Version
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FILING DATE:

11-NOV-1996

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Patent No. 6521737
GENERAL INFORMATION:
APPLICANT: O'Mahony, Daniel J
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                              TELEFAX: 770 534-8247
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
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DESCRIPTION:
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ATTORNEY/AGENT INFORMATION:
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REFERENCE/DOCKET NUMBER: 97.1061.US
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                                                                                                 REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/746,411
FILING DATE: 08-NOV-1996
APPLICATION NUMBER: US 60/006461
FILING DATE: 10-NOV-1995
APPLICATION NUMBER: IE 950865
FILING DATE: 10-NOV-1995
                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                    ATTORNEY/AGENT INFORMATION: NAME: Severson, Mary L
                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
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FILING DATE: 19-Aug-2002
CLASSIFICATION: «Unknown»
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 TYPE: nucleic
                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 1300 Gould CITY: Gainesville
LENGTH: 126 base pairs
TYPE: nucleic acid
                                                                                 TELEPHONE: 770 534-8239
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/desc = "individual isolate"
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Tissues and Methods of Identifying and Using the Same
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                                                                                                                                      34,927
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                                                              Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Patent No. 561247
                                                                                                                                                                                                                                                                                                                                      TELEFAX: (415) 543-5043
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/248,474
FILING DATE: 25-MAY-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                           FEATURE:
                                                                                                                                                                                                                                        MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: MCK. BIRD, Davi
APPLICANT: WILSON, Mark A.
                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Bastian, Kevin L.
REGISTRATION NUMBER: 34,774
REFERENCE/DOCKET NUMBER: 23
TELECOMMUNICATION INFORMATION:
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ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                          NAME/KEY: misc_feature
LOCATION: 1..243
                                                                                                                                                                                                          ORGANISM:
                                                                                                                                                                                                                                                                        TOPOLOGY:
                                                                                                                                                                                                                                                                                      STRANDEDNESS:
                                                                                                                                                                                                                                                                                                LENGTH: 243 base pairs
TYPE: nucleic acid
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MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "individual isolate"
SEQUENCE DESCRIPTION: SEQ ID NO: 9:
133 TTAGGAGGAACAA 121
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                                                                              Similarity
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                                                            65.0%; Score 13; ilarity 100.0%; Pred. No. Conservative 0; Mismatc
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100.0%; Pred. No.
:ive 0; Mismatc
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                                                               Mismatches
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76;
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                                                                                             Length 243;
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RESULT 27

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RESULT 28
US-09-702-705-1692/c
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Best Local :
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                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                            Sequence 1692, Apparent No. 65040
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (415) 576-0300 INFORMATION FOR SEQ ID NO: 23: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION: David McK.
APPLICANT: Wilson, Mark A.
APPLICANT: Wilson, Mark A.
TITLE OF INVENTION: Nematode
                                                                APPLICANT:
                                                                                            APPLICANT:
                                                                                                             APPLICANT:
                                               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Bastian, Kevin L.
REGISTRATION NUMBER: 34,774
REFERENCE/DOCKET NUMBER: 02
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 06
FILING DATE: 25-WAY-1994
ATTORNEY/AGENT INFORMATION:
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ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: 1
                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: misc_feature
LOCATION: 1.243
OTHER INFORMATION: /standard_name= "DB# 139"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: FILING DATE: 26-NOV
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CITY: San Francisco
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 243 base pairs
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                                                                                                                                                                                                                                                         133 TTÁGGÁGGÁÁCAÁ 121
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          Carter, Darrick
Retter, Marc
Mannion, Jane
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Fan, Liqun
                                                                                           Bangur, Chaitanya
Lodes, Michael A.
                                                           Vedvick, Tom
                                                                          Fanger, Gary
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                                                                                                                                                                          Application US/09702705
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                                                                                                             Chaitanya
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; Pred. No.
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76;
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RESULT 30 US-08-244-205-4/c

Sequence 4, Application US/08244205 Patent No. 5952544 GENERAL INFORMATION:

APPLICANT:

Browse, John, Kinney, Anthony J.,

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FILE REFERENCE: 210121.478C14

FILE REFERENCE: 210121.478C14

CURRENT APPLICATION NUMBER: US/09/702,705

CURRENT FILING DATE: 2000-10-30

NUMBER OF SEQ ID NOS: 1833

SOFTWARE: FASTSEQ for Wird

SEQ ID NO 1692

LENGTH: 15
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US-09-736-457-1692/c
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; OTHER INFORMATION: n = A,T,C or US-09-702-705-1692
                                                                                                        US-09-736-457-1692
                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/736,457
CURRENT FILING DATE: 2000-12-13
NUMBER OF SEQ ID NOS: 1864
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 1662
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                    Matches
                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent No. 650944
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc feature
                                                                Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                 APPLICANT: Wang, Aijun
TITLE OF INVENTION: COMPOSITIONS AND METHODS
TITLE OF INVENTION: DIAGNOSIS OF LUNG CANCER
FILE REFERENCE: 210121.478C15
                                                                                                                  NAME/KEY: misc_feature
LOCATION: (1)...(450)
OTHER INFORMATION: n = A,T,C or
                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                            FEATURE:
                                                                                                                                                                       ORGANISM: Homo sapiens
                                                                                                                                                                                     LENGTH: 450
TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION:
235
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                       2 CTTAGGAGGAACA 14
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CTTAGGAGGAACA 223
                                                                                                                                                                                                                                                                                                                                     Fanger, Gary
Vedvick, Tom
Carter, Darrick
Retter, Marc
Mannion, Jane
Fan, Ligun
                                                                                                                                                                                                                                                                                                                                                                                                                      Wang, Tongtong
Bangur, Chaitanya
Lodes, Michael A.
                                                    Conservative
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100.0%; Prr
                                                               65.0%;
                                                               Score 13;
; Pred. No.
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Pred. No.
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                                                     Mismatches
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                                                                DB 4;
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73;
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COUNTRY: U.S.A.

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LOCATION:
US-08-244-205-4
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                                                                                                                                                                                                                RESULT 31
                                                                                                                                                               Sequence 4, Application PC/TUS9210284 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 07/804,259
FILLING DATE: 4 DECEMBER 1991
ATTORNEY/AGENT INFORMATION:
NAME: Floyd, Linda A.
REGISTRATION NUMBER: 33,692
REFERENCE/DOCKET NUMBER: BB-10
TELECOMMUNICATION INFORMATION:
TELEPHONE: (302) 992-4929
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (302) 892-794
TELEX: 835420
NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                              APPLICANT: Browse, John, Kinney, Anthony J., APPLICANT: Pierce, John, Wierzbicki, Anna M., APPLICANT: Yadav, Narendra S., Perez-Grau, Luis TITLE OF INVENTION: Fatty Acid Desaturase Genes TITLE OF INVENTION: from Plants
                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLONE: PACF2-2 FEATURE:
                                              NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           UKGANISM: Arabidopsis thaliana IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORIGINAL SOURCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: Macintosh
OPERATING SYSTEM: Macintosh System,
SOFTWARE: Microsoft Word, 4.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: 9 US/08/244,205
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: cI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (302) 892-7949
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: 1007 Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 1525 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: 800
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                 STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: E. I. du Pont de Nemours and Company
                              ADDRESSEE:
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Wilmington
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             E: E. I. du Pont de
1007 Market Street
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                                I. du Pont de Nemours and Company
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Fatty Acid Desaturase Genes
from Plants
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                                                                                                                                                                                                                                                                                                                                                   DB 2;
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PCT-US94-01321-9/c
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FILING DATE: 05-FEB-1993
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/156551
FILING DATE: 22-NOV-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/014431
                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIOM TYPE: Plopy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/01321
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Microsoft Word, 4.0
CURRENT APPLICATION DATA:
APPLICATION NITHINGS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
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REFERENCE/DOCKET NUMBER: BE
TELECOMMUNICATION INFORMATION:
TELEPHONE: (302) 992-4929
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 07/804,2
FILING DATE: 4 DECEMBER 1991
ATTORNEY/AGENT INFORMATION:
NAME: Floyd, Linda A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                    TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: CI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 1525 base pairs
TYPE: NUCLEIC ACID
STRANDEDNESS: single
                                                                                                                                    CLASSIFICATION:
                                                                                                                                                      FILING DATE: . 04-FEB-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LOCATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            814 GGAGGAACAAGTC 802
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                                                                                                                                                                                                                                                                                      Altered Linolenic and Linoleic Acid Content in Plants
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1645 base pairs

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; TYPE: DNA
; ORGANISM: homo sapiens
US-10-020-079-39
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US-10-020-079-39/c
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US-09-389-956-79/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; FEATURE:
; NAME/KEY: CDS
; LOCATION: (1)..(1596)
US-09-389-956-79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCT-US94-01321-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Huang, Shi
TITLE OF INVENTION: PR.Domain Containing Nucleic Acids, Polypeptides,
TITLE OF INVENTION: Antibodies and Methods
FILE REFERENCE: P-LJ 3611
CURRENT APPLICATION NUMBER: US/09/389,956
CURRENT FILING DATE: 1999-09-03
NUMBER OF SEQ ID NOS: 93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 79, Application US/09389956 Patent No. 6586579
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 65.0%;
Best Local Similarity 100.0%;
Matches 13; Conservative (
                                                                         SEQ ID NO 39
                                                                                                                                                                                                                                                                                                                                          Sequence 39, Application US/10020079
Patent No. 6579710
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                           APPLICANT: Turner, C. Alexander Jr.
APPLICANT: Mathur, Brian
APPLICANT: Friddle, Carl Johan
TITLE OF INVENTION: No. 6579710el Human Kinases and Polynucleotides Encoding the Same
FILE REFERENCE: LEX-0281-USA
CURRENT APPLICATION NUMBER: US/10/020,079
CURRENT FILING DATE: 2001-12-12
                                                                                                                        PRIOR FILING DATE: 2000-12-12
PRIOR APPLICATION NUMBER: US 60/289,422
PRIOR FILING DATE: 2001-05-08
                                                                                                                                                                                                                                                                                                                           ENERAL INFORMATION:
                                                                                      NUMBER OF SEQ ID NOS: 40
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                              PRIOR APPLICATION NUMBER: US 60/255,103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 1970
TYPE: DNA
ORGANISM: Homo sapiens
                                                    LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: cDNA
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LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                      139 AGGAGGAACAAGT 127
                                                      2517
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CURRENT APPLICATION NUMBER: US/10/020,079
CURRENT FILING DATE: 2001-12-12
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This primer binds to a region at the 5' end of the Epstein-Barr virus
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                                                                                                                                                                                                 New compositions comprising nucleic acid sequences which hybridizes to Epstein-Barr virus (EBV) nucleic acid, for in clinical specimens to determine patients at high risk
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                                                                                                                                                                            EBV infections
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2002WO-US04339.
                                                                                                                          44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR;
                                                                                                                                                                                                                                                                                                                                                         PA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             probe; detection; Epstein-Barr virus;
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                                                                                                                          English.
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Pred. No.
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The sequences given in ABA00268-75 are primers and probes which used in the compositions of the invention for the detection of Epstein-Barr virus (EBV). The compositions comprise at least or purified and isolated oligonucleotide consisting of a nucleic ac

acid

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cc are useful for detecting EBV in clinical specimens to determine cc patients who are at high risk to develop serious and compliations comprising them ccomplications, and allow for better clinical management of these patients by earlier recognition of their infection status. The cc oligonucleotide sequences may also be used to amplify EBV DNA cc detecting EBV has a broad dynamic range of detection from less than cc 10-10000000 copies of EBV DNA, is less labour intensive requiring only one reaction tube for the EBV determination, highly sensitive, accurate and has a rapid turn around time with assays that are completed, including amplification, probe specific hybridization, and calculation of copy number in less than 1 hour. The method may be adapted to c automated systems.
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Best Local
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                                                                                New fragment of Epstein-Barr Virus DNA - useful in vector to express polypeptide for use in prepn. of vaccine against the virus and for use in diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-MAR-2003
17-OCT-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
The sequence encodes an outer surface viral protein of EBV, used to generate antibodies reacting with the surface proteins of EBV-infected cells, and in the preparation of a vaccine against I (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                     Kleff E,
                                                                                                                                                                                                                      23-JUL-1984;
30-JAN-1984;
                                                                                                                                                                                                                                                           28-JAN-1985;
                                                                                                                                                                                                                                                                                                           EP151079-A.
                                                                                                                                                                                                                                                                                                                                                                                                Epstein-Barr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAN50114 standard; DNA;
                                                         Claim 1;
                                                                                                                                 P-PSDB;
                                                                                                                                              WPI; 1985-191978/32.
                                                                                                                                                                                               (UYCH-) UNIV CHICAGO.
                                                                                                                                                                                                                                                                                    07-AUG-1985
                                                                                                                                                                                                                                                                                                                                                          mat_peptide
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                                                           Page 21-23;
                                                                                                                                                                      Tanner J,
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                                                                                                                                                                                                                                                                                                                                                                                                                       virus; antigen; vaccine; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                              encoding Epstein-Barr virus (EBV)
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(first entry)
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84US-0575352.
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                                                                                                                                                                                                                                                                                                                              /*tag= a
/label= EBV surface protein antigen
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G;
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Pred. No. 0.078;
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           against EBV
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       The donor and acceptor splice sites of the EBV gene encoding gp350/ 220 are mutated by replacement of native nucleotides by non-native nucleotides, without altering the encoded amino acid sequence, resulting in elimination of gp220 prodn. Recombinant homogeneous gp350, useful in vaccines, is expressed in mammalian or insect cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18-JAN-1996
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                                                                                         New DNA encoding a homogeneous gp350 protein - can preventing and treating Epstein-Barr virus-related
                                                                                                                                                                                                                                                                                                                     misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                     sig_peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Epstein-Barr virus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EBV gp350/220 cDNA
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                                                              Claim 2; Fig.1; 61pp;
                                                                                                                                                 Jackman WT,
                                                                                                                                                                                     18-APR-1994;
                                                                                                                                                                                                       13-APR-1995;
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                                                                                conditions
                                                                                                                     P-PSDB; AAR80144.
                                                                                                                                                                   (AVIR-) AVIRON
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Conservative
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/*raq= f
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1014..3737
1*tag= a
1014..1067
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/*tam-
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                                                                                                                                                                                                                                                                                                                               (*tag= d splice donor site /function= splice donor site /function= "bases 2513-2517 (AAGT)
                                                                                                                                                                                                                                                                               function= splice acceptor site /note= "bases 3104-3107 (AGGT) ar TGGA in the non-splicing
                                                                                                                                                                                                                                                                                                            *tag=
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Pred.
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The invention relates to an isolated polynucleotide (I) comprising: (a) CC any of a group of over 4000 nucleotide sequences (ABV94628-ABV99145); CC (b) complements of (a); (c) sequences consisting of at least 20 contiguous residues of (a); (d) sequences that hybridize to (a), under moderately stringent conditions; (e) sequences having at least 75% or 90% (C (ABP68596-ABP68637) encoded by (I) and oligonucleotide can be used to detect cancer in a patient and compositions comprising polypeptides (C antigen presenting cells expressing the polypeptide are useful in CC antigen presenting cells expressing the polypeptide are useful in CC treating pancreatic cancer and stimulating an immune response. The polynucleotides can be used as grobes or primers for nucleic acid (c) hybridisation, in the design and preparation of ribozyme molecules for inhibiting expression of the tumour polypeptides and proteins in the cumour cells, in vaccines and for gene therapy.

CC Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at figure 1.
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Best Local S
Matches 20
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                                                                                                                                                                                                                                                                                                                                                                                               Benson
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28-APR-2001;
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31-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
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                                                                                                                                                                                                                                                                                                       isolated polynucleotide and pancreatic tumor polypeptides, useful diagnosing, preventing and/or treating cancer, particularly creatic cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2899
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; 2001US-265682P.
; 2001US-267568P.
; 2001US-278651P.
; 2001US-287112P.
; 2001US-291631P.
; 2001US-3305484P.
; 2001US-313999P.
; 2001US-333626P.
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Pred. No. 0.055;
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Best Local
                                                 identity to (a); or (f) degenerate variants of (a). Polypeptides (ABP68596-ABP68637) encoded by (I) and oligonucleotide can be used to detect cancer in a patient and compositions comprising polypeptides, polynucleotides, antibodies, fusion proteins, T cell populations and antigen presenting cells expressing the polypeptide are useful in treating pancreatic cancer and stimulating an immune response. The polynucleotides can be used as probes or primers for nucleic acid hybridisation, in the design and preparation of ribozyme molecules for inhibiting expression of the tumour polypeptides and proteins in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-MAY-2001;
12-JUL-2001;
20-AUG-2001;
          tumour cells, in vaccines and for gene therapy.

Note: The sequence data for this patent did not form
specification, but was obtained in electronic format
                                                                                                                                                                 The invention relates to an isolated polynucleotide (I) comprising: (a) any of a group of over 4000 nucleotide sequences (ABV94628-ABV99145); (b) complements of (a); (c) sequences stating of at least 20 contiguous residues of (a); (d) sequences that hybridize to (a), under moderately stringent conditions; (e) sequences having at least 75% or 9 identity to (a); or (f) degenerate variants of (a). Polypeptides
                                                                                                                                                                                                                                                                                             New isolated polynucleotide and pancreatic tumor polypeptides, useful for diagnosing, preventing and/or treating cancer, particularly pancreatic cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; pancreas; cancer; gene therapy; vaccine; immunostimulant;
cytostatic; tumour; gene; ss.
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                                                                                                                                                                                                                                                                    Claim 1;
                                                                                                                                                                                                                                                                                                                                                      WPI; 2002-627435/67.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-JAN-2003
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                                                                                                                                                                                                                                                                    SEQ
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                                                                                                                                                                                                                                                                                                                                                                                Kalos MD,
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2001US-265682P.
2001US-267568P.
2001US-278651P.
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2001US-313999P.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                  Hepler WT,
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              part of the printed directly from WIPO
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                                                              Query Match
Best Local
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                                                                                                                   monitoring viral gene expression or activity during a treatment regime. The nucleic acid molecules are also useful as antisense constructs to control viral gene expression in infected cells and tissues and to cretransgenic viral resistant shrimp.
                                                                                                                                                         The invention provides the primary nucleotide sequence of the WSBV genome (AAH62689), predicted transcript sequences (AAH62689-AAH62839) and encoded proteins (AAG684910-AAG685051) and oligonucleotide sequences (AAH62840-63160) suitable for use as primers or probes. The nucleic acid molecules and proteins of the invention are useful for diagnosis and monitoring viral infection, in screens for antiviral agents and for monitoring viral gene expression or activity during a treatment regimen.
                                                                                                                                                                                                                                                                                   Primary nucleotide sequence of the shrimp white spot Bacilliform virus (WSBV), useful for producing viral polypeptides that can be used to screen for agents that are useful for treating WSBV infection -
                                                                                                                                                                                                                                                                                                                                                                          Xu X,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Shrimp white spot Bacilliform virus; WSBV; diagnosis; viral infection; antiviral agent; gene expression; antisense construct; transgenic viral resistant shrimp; ds.
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                                                                                               Sequence 2376 BP; 787 A; 467 C; 526 G;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    White spot syndrome virus.
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                                                                                                                                                                                                                                                             Claim 4; Figure 3; 626pp; English
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DB; AAG84928.
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16; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                  THIRD INST OCEANOGRAPHY STATE OCEANI C SINOGENOMAX CO LTD.
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                                                            Similarity
                  AGGAGGAACAAGTCCC 20
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                                                 Conservative
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                                                        80.0%; Score 16;
100.0%; Pred. No.
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Pred. No.
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                                                                                               596 T; 0 other;
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                                                                       Length 2376;
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CTTAGGAGGAACAAGT

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80.0%; llarity 100.0%; Conservative

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8. BB . 22; Sequence 6572

BP; 1929 A; 1256 C; 1418 G; 1969

T; 0 other; Length 6572;

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Zhao QA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Alzheimer's; Parkinson's disease; Huntington's disease; amyotrophic lateral sclerosis; Shy-Drager Syndrome; chemchemokinetic; thrombolytic; drug screening.
                                                                                                                      system, such as peripheral nervous injuries, peripheral neuropathy and localised neuropathies and central nervous system diseases, such as Alzheimer's, Parkinson's disease, Huntington's disease, amyotrophic lateral sclerosis, and Shy-Drager Syndrome. Other uses include the utilisation of the activities such as: Immune system suppression, Activin/inhibin activity, chemotactic/chemokinetic activity, haemostatic and thrombolytic activity, cancer diagnosis and therapy, drug screening, assays for receptor activity, arthritis and inflammation, leukaemias and
                                                                                                                                                                                                                                                                                                                                 immunosuppressant and cytostatic activity. The polynucleotides are useful in gene therapy. A composition containing a polypeptide or polynucleotide of the invention may be used to treat diseases of the peripheral nervous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel nucleic acids and polypeptides, useful such as central nervous system injuries -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21-JAN-2000;
25-APR-2000;
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                                                                        C.N.S disorders.
Note: The sequence
                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to human nucleic acids (AAI57798-AAI61369) and the encoded polypeptides (AAM38642-AAM42213) with nootropic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-442253/47.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-JUL-2000;
03-AUG-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-JUL-2000;
                                                  specification.
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Wang Z,
Zhou P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 1134;
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2000US-0727344
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2000US-0598042.
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2000US-0653450.
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Wehrman T, X
Goodrich R,
                                                                           data
                                                                           for this patent did not form part
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10078pp; English.
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Xu C, Xue AJ,
R, Drmanac RT;
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ervous system; CNS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        treating
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ren F,
Zhang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     inflammation;
                                                                           of the printed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Wang
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В

4524

CTTAGGAGGAACAAGT 4539

ABA09554 standard; cDNA; 7445 BP

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ABA09554
ABA0954
ABA09554
ABA09554
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ABA09554
ABA09554
ABA09554
ABA0954
ABA09554
ABA0964
ABA09
                                                                                                                                                                                                                                                                   Sequences ABB10981-ABB12330 represent 1350 novel human polypeptides, and converted to the first provided to the first polypeptides and recombinant host cells comprising a nucleotide of the invention, methods of producing the novel polypeptides, antibodies against the polypeptides, methods of detecting the nucleotides or polypeptides in a sample, and methods of identifying compounds which bind to polypeptides of the invention. Although novel, many of the complete of the invention although novel, many of the compounds which bind to polypeptides of the invention have homology to known proteins, thereby giving an insight into their probable biological activities, and hence contential therapeutic applications. The polypeptides of the invention may have various activities, including cytokine, cell proliferation or cell differential therapeutic applications. The polypeptides of the invention may have various activities; stem cell growth activity; have various activity; activity or inhibin-related activities; themostatic or chemotactic or chemokinetic activities, hamostatic, thrombotic or chemokinetic activities, hamostatic, thrombotic or
conditions, e.g., by protein or gene therapy. Such conditions include cancers, haematopoietic disorders (e.g., myeloid or lymphoid cell disorders), chronic inflammatory conditions (e.g., asthma or arthritis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; cytokine; cell proliferation; cell differentiation; growth factor; haematopoiesis regulation; tissue growth; immunomodulator; activin; inhibin; chemotaxis; chemokinesis; thrombolysis; oncogenesis; proliferation; metastasis; cancer; tumour; haematopoietic disorder; myeloid cell disorder; lymphoid cell disorder; asthma; arthritis; chronic inflammatory condition; proliferative retinopathy; atherosclerosis; coronary heart disease; arterial ischaemia;
                                                                                                                 thrombolytic activities; receptor or ligand activities; or may be involved in oncogenesis, cancer cell proliferation or metastasis. Depending on their biological activities, polypeptides and nucleotides of the invention are useful for preventing, treating or ameliorating medical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 992-994; 1963pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        e.g. arthritis and cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human proteins and DNA encoding sequences useful for preventing, treating or ameliorating a medical condition in a mammalian subject
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-457740/49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-FEB-2000;
27-APR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-FEB-2001; 2001WO-US03800.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200157188-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cytostatic; osteopathic; vasotropic; cardiant; virucide; antibacterial; antifungal; vulnerary; antiulcer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cell culture; drug screening; gene therapy; antiinflammatory;
antiasthmatic; antiarthritic; haemostatic; antiarteriosclerotic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                bone disorder; osteoporosis; vascular growth disorder; tissue regeneration; wound healing; infection; immune disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human guanine nucleotide exchange factor homologue cDNA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (HYSE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ŢŢ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABB12310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Liu C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2000US-0496914.
2000US-0560875.
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            asthma or arthritis),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO:1330.
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RESULT 10
AAK51763
ID AAK51763
XX AAK517
AC AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CC proliferative retinopathy, atherosclerosis, coronary heart disease,
CC arterial ischaemia, bone disorders (e.g., osteoporosis), and abnormal
CC vascular growth. Polypeptides involved with tissue regeneration and
CC repair (or nucleic acids encoding them) may be used to promote wound
CC healing (e.g., of burns, incisions and ulcers), while those with
CC immunomodulatory activities may be used in the treatment of viral,
CC bacterial and fungal infections in addition to immune disorders.
CC Polypeptides with growth factor activity may be used in cell cultures to
CC promote cell growth. For example, such polypeptides may be used to
CC manipulate stem cells in culture to give rise to neuroepithelial cells
CC that can be used to augment or replace cells damaged by illness,
CC autoimmune disease or accidental damage. The polypeptides and nucleotides
CC may also be used in the diagnosis of the above conditions, and in drug
CC screening techniques. The present sequence represents a cDNA encoding a
CC novel human polypeptide of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-FEB-2000; 2
27-APR-2000; 2
20-JUN-2000; 2
19-JUL-2000; 2
01-SEP-2000; 2
                                                                                                                                                                                                                                                               Tang YT,
Zhao QA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human, cytokine, cell proliferation, cell differentiation, gene therapy, vaccine, peptide therapy, stem cell growth factor, haematopoiesis, tissue growth factor; immunomodulatory, cancer; leukaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human polynucleotide SEQ ID NO 308
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAK51763 standard; cDNA; 7445 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                           WPI; 2001-476283/51.
P-PSDB; AAM78630.
                                                                                                                                                                                                                                                                                                                                              (HYSE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-FEB-2001; 2001WO-US04098
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nervous system disorder; arthritis; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                          20-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   L5-SEP-2000;
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                                                                                                                                                                                                                                                                                                                                              HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 7445
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                                                                                                                                                                                                                                 Liu C,
Wang D,
Yang Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CTTAGGAGGAACAAGT
                                                                                                                                                                                                                                                                                                                                                                                            ; 2000US-0663561.
; 2000US-0693325.
; 2000US-0728422.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2000US-0496914.
2000US-0560875.
2000US-0598075.
2000US-0620325.
2000US-0654936.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 2177 A; 1404 C; 1582 G; 2282
                                                                                                                                                                                                                                    Wang J, Zi
Wejhrman T,
                                                                                                                                                                                                                                                                                           Drmanac
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     80.0%;
                                                                                                                                                                                                                                    RT, Asundi V, Zhou
, Zhang J, Ren F, (
n T, Goodrich R;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 16; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 22;
. 8.7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 7445;
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                                                                                                                                                                                                                                                                     u C,
Wang,
                                                                                                                                                                                                                                                  Cao .
'g Zw;
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Claim 1; Page 1279-1284; 6221pp; English

useful in diagnosis and

gene

therapy

Nucleic acids encoding polypeptides with cytokine-like activities

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RESULT 11
AAK5277
ID AAK52
XX AAK52
XX AAK52
XX O6-NO
DT O6-NO
DE Human
XX Human
XX Human
XX WO200
XX WO200
XX WO200
XX WO200
XX O9-AU
PF 05-FE
XX 03-FE
PR 27-AE
PR 27-AE
PR 19-JU
PR 19-FSI
XX WPI
PR 20-G
PI VUE
PR 20-G
PI VUE
PR 20-WE
PR 20-WE
PR 20-WE
PR 20-WE
PR 30-WE
PR 20-WE
P
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-FEB-2000; 2000US-0496914.

27-APR-2000; 2000US-0560875.

20-JUN-2000; 2000US-0598075.

19-UUL-2000; 2000US-0620325.

01-SEP-2000; 2000US-0654936.

15-SEP-2000; 2000US-0663561.

20-OCT-2000; 2000US-0693325.

30-NOV-2000; 2000US-0728422.
                                                                                                                                                                                                                                                                                                                                                                                                                   Tang YT,
Zhao QA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to polynucleotides (AAK51456-AAK53435) and the encoded polypeptides (AAM78323-AAM80302) that exhibit activity elating to cytokine, cell proliferation or cell differentiation or which may induce production of other cytokines in other cell populations. The polynucleotides and polypeptides are useful in gene therapy, vaccines or peptide therapy. The polypeptides have various cytokine-like activities, e.g. stem cell growth factor activity, haemaropoiesis regulating activity, tissue growth factor activity, immunomodulatory activity and activiny inhibit activity and may be useful in the diagnosis and/or restrict the control of 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; cytokine; cell_proliferation; cell differentiation; gene therapy; vaccine; peptide therapy; seem cell growth factor; haematopoiesis; tissue growth factor; immunomodulatory; cancer; leukaemia; nervous system disorder; arthritis; inflammation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         inflammation.

Note: Records for SEQ ID NO 2110 (AAK52581), 2111 (AAK52582) and 3666 (AAM80020) are omitted as the relevant pages from the sequence listing
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                                                                                                                                                    Nucleic acids encoding polypeptides with cytokine-like useful in diagnosis and gene therapy -
                                                                                                                                                                                                                                                                  WPI; 2001-476283/51.
P-PSDB; AAM79614.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human polynucleotide SEQ ID NO 2276
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAK52747 standard; cDNA; 7445 BP
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                                                                          Claim 1; Page 4614-4616; 6221pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200157190-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16;
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                                                                                                                                                                                                                                                                                                                                                                                Yang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CTTAGGAGGAACAAGT 4496
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                                                                                                                                                                                                                                                                                                                                                                            Liu C,
Wang D,
Yang Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                            Drmanac RT, Asundi V,
Wang J, Zhang J, Ren
Wejhrman T, Goodrich R;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 2;
8.7;
                                                                                                                                                                                                                                                                                                                                                                                                               Zhou
F, C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22;
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                                                                                                                                                                                                                                                                                                                                                                                                                       Chen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                   Xu C, Ca
                                                                                                                                                                                               activities
                                                                                                                                                                                                                                                                                                                                                                                                                       Cao Y,
ng ZW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ma
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The invention relates to

polynucleotides (AAK51456-AAK53435) and the

Claim 1;

SEQ ID NO 4706; 10078pp; English

such as

central nervous

polypeptides, useful system injuries -

for treating disorders

Novel nucleic acids and

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RESULT 12
AAI60717
ID AAI60717
AAI60717
AAI60717
AC AAI60
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                               21-JAN-2000; 2000US-0488725.

25-APR-2000; 2000US-0552317.

09-JUL-2000; 2000US-0598042.

19-JUL-2000; 2000US-0620312.

03-AUG-2000; 2000US-0653450.

14-SEP-2000; 2000US-0662191.

19-OCT-2000; 2000US-0693036.

29-NOV-2000; 2000US-0727344.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; nootropic; immunosuppressant; cytostatic; gene therapy; cancer; peripheral nervous system; neuropathy; central nervous system; CNS; Alzheimer's, Parkinson's disease, Huntington's disease; haemostatic; amyotrophic lateral sclerosis; Shy-Drager Syndrome; chemotactic; chemokinetic; thrombolytic; drug screening; arthritis; inflammation;
                                                                                                                                                                                                                             Tang
Wang
Zhao
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human polynucleotide SEQ ID NO 4706.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 7445 BP; 2177 A; 1404 C; 1582 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Note: Records for SEQ ID NO 2110 (AAK52581), (AAM80020) are omitted as the relevant pages were missing at the time of publication.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AA160717 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       treatment of cancer, leukaemia, nervous system disorders, arthritis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26-DEC-2000; 2000WO-US34263
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200153312-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                     WPI; 2001-442253/47.
P-PSDB; AAM41561.
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                                                                                                                                                                                                                                                                                                                                  (HYSE-) HYSEQ INC.
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                                                                                                                                                                                                                             Š, Ž
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               l Similarity
16; Conserv
                                                                                                                                                                                                                          Liu C,
Wang Z,
Zhou P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CTTAGGAGGAACAAGT 4507
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                                                                                                                                                                                                                             Asundi V,
Wehrman T,
Goodrich F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 80.0%;
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Pred. No.
                                                                                                                                                                                                                                                       Chen R,
Xu C,
                                                                                                                                                                                                                          hen R, Ma Y,
Ku C, Xue AJ,
Drmanac RT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2282
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from
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Yang
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                                                                                                                                                                                                                                                       AX P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0 other;
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                                                                                                                                                                                                                                                          ŗ,
                                                                                                                                                                                                                                                                               Wang
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RESULT 13
AAS82621
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           immunosuppressant and cytostatic activity. The polynucleotides are useful in gene therapy. A composition containing a polypeptide or polynucleotide of the invention may be used to treat diseases of the peripheral nervous system, such as peripheral nervous injuries, peripheral neuropathy and localised neuropathies and central nervous system diseases, such as Alzheimer's, Parkinson's disease, Huntington's disease, amyotrophic lateral sclerosis, and Shy-Drager Syndrome. Other uses include the utilisation of the activity, chemotactic/chemokinetic activity, haemostatic and thrombolytic activity, cancer diagnosis and therapy, drug screening, assays for receptor activity, arthritis and inflammation, leukaemias and
polypeptide (II) sequences. (I) is useful as hybridisation probes, polymerase chain reaction (PCR) primers, oligomers, and for chromosome polymene mapping, and in recombinant production of (II). The polymers polymerase are also used in diagnostics as expressed sequence tags for identifying expressed genes. (I) is useful in gene therapy technique to restore normal activity of (II) or to treat disease states involving
                                                                                                                                                                             New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess
                                                                                                     The
                                                                                                                                 Claim 1; SEQ ID No 18425; 103pp;
                                                                                                                                                                 biodiversity
                                                                                                                                                                                                                                                p-PSDB;
                                                                                                                                                                                                                                                                                                                                                              31-MAR-2000;
23-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                             30-MAR-2001; 2001WO-US08631
                                                                                                                                                                                                                                                                                                                                                                                                                                             11-OCT-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200175067-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-FEB-2002
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                                                                                                 invention relates to isolated polynucleotide (I) and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4492
                                                                                                                                                                                                                                                                 2001-639362/73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          supplement;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        chromosome mapping; gene mapping; gene therapy;
upplement; medical imaging; diagnostic; genetic of
upplement; medical imaging;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 Similarity
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                                                                                                                                                                                                                                                                                                 RT,
                                                                                                                                                                                                                                                ABG18434.
                                                                                                                                                                                                                                                                                                                              HYSEQ
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llarity 100.0%;
Conservative 0;
                                                                                                                                                                                                                                                                                              Liu C,
                                                                                                                                                                                                                                                                                                                                                            2000US-0540217.
2000US-0649167.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         novel human diagnostic protein #18425.
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                                                                                                                                                                                                                                                                                              Tang
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Mismatches
                                                                                                                                English.
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8.7;
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RESULT 14
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JID ACC500
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PR 25-5E
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PR 18-JU
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (II). (II) is useful for generating antibodies against it, detecting or quantitating a polypeptide in tissue, as molecular weight markers and as a food supplement. (II) and its binding partners are useful in medical imaging of sites expressing (II). (I) and (II) are useful for treating disorders involving aberrant protein expression or biological activity. The polypeptide and polynucleotide sequences have applications in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits to assess biodiversity and to produce other types of data and products dependent on DNA and amino acid sequences. AAS64197-AAS9454 represent novel human diagnostic coding sequences of the invention.

Note: The sequence data for this patent did not appear in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                  Lillie J, (
                                                                                                                                                                                     Breast cancer diagnosis or treatment expression of a marker in a patient snon-breast cancer sample -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-JUN-2001;
27-JUN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 7843 BP; 2262 A; 1492 C; 1682 G; 2403 T; 4 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO2003004989-A2
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                                                                                                                                           Claim 1;
                                                                                                                                                                                                                                                                                         P-PSDB;
                                                                                                                                                                                                                                                                                                                                                         Hortobagyi GN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-MAR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18-JUL-2001;
25-SEP-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; breast cancer; cytostatic; gene therapy; gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Breast cancer associated cDNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                             (MILL-) MILLENIUM PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14-MAY-2002;
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                                                                                                                                                                                                                                                                                         ABR47392
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                                                                                                                                           SEQ ID 14; 128pp;
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                                                                                                                                                                                                                                                                                                                                                                                     Gannavarapu M,
, Myer V, Wang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2001US-306501P.
2001US-325002P.
2002US-362585P.
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2001US-301572P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2002US-380391P
                                                                                                                                                                                                                                                                                                                                                              Pusztai L,
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                                                                                                                                                                                                                                                                                                                                                              Glatt K,
y Y, Xu Y,
Meric F,
                                                                                                                                           English.
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Pred. No.
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                                                                                                                                                                                                                 sample with that i
                                                                                                                                                                                                                                                                                                                                                              Hoersh S,
Zhao X,
Sahin A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO:14.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23;
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                                                                                                                                                                                                                                                                                                                                                                   Meyers
Mills
                                                                                                                                                                                                                                                                                                                                                                 Kamatkar
Meyers RE,
Mills GB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length
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                                                                                                                                                                                                                                                                                                                                                                                                              Mertens M;
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The present invention describes a method for assessing whether a patient is afflicted with breat cancer. The method comprise comparing the level of expression of a marker (gene/polypeptide see ACC50076 to ACC50334 and ABR47386 to ABR47632) in a patient sample and the normal level of

sample,

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Matches
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The invention relates to human novel polynucleotides and associated polypeptides. The polynucleotides and polypeptides are useful for treating inflammatory conditions such as arthritis, nephritis, Crohn's disease, ischaemia-reperfusion injury, shock, sepsis, immune responses and cancer and for promoting wound healing. The sequences are used to induce the proliferation of neural cells and regeneration of nerve and brain tissue, and are useful for the treatment of central and peripheral nervous system diseases and neuropathies, such as Alzheimer's disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; gene; ss; inflammatory condition; shock; sepsis; immune response; cancer; wound healing; central nervous system disease; haematopoiesis; peripheral nervous system disease; atteral sclerosis; tendon; myeloid cell disorder; lymphoid cell disorder; platelet disorder; bone; cartilage; ligament; nerve tissue; ulcer; osteoporosis; osteoparthritis; bone degenerative disorder; periodontal disease; reperfusion injury; lung fibrosis; liver fibrosis; autoimmune disorder; bacterial infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                patient sample and the normal level is an indication that the patient afflicted with breast cancer. The breast cancer associated sequences from the present invention have cyrostatic activities and can be used gene therapy. The method is useful for diagnosing and treating breast
                                                                                                                                                                                                                                                                                                                      Novel nucleic acids and polypeptides for diagnosis, inflammatory, autoimmune, nervous system, myeloid o disorders, cancer and promoting wound healing -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-AUG-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tang YT, G
Yamazaki V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-NOV-2001; 2001WO-US47004.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human novel polynúcleotide #40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABK94929 standard; cDNA; 9783 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-508509/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-NOV-2000; 2000US-0028952
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           allergic
                                                                                                                                                                                                                                                                  Claim 1; Page 425-434; 672pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (HYSE-) HYSEQ INC
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16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           condition; thrombolysis; thrombosis; coagulation disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CTTAGGAGGAACAAGT
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V, Ujwal
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Drmanac
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Pred. No.
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RT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2735
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RESULT 16
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16-MAR-2000; 2000US-189862P.
25-MAY-2000; 2000US-207454P.
09-JUN-2000; 2000US-211314P.
18-JUL-2000; 2000US-219907P.
13-DEC-2000; 2000US-255281P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            reperfusion injury in various tissues, immune deficiencies and disorders including severe combined immunodeficiency (SCID), bacterial or fungal infections, autoimmune disorders [50] multiple sclerosis and myasthenia gravis, allergic conditions such as asthma, thrombolysis or thrombosis and coagulation disorders. Sequences ABK94890-ABK94982 represent human novel polynucleotides of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-SEP-2002
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                                                                                                                                                                                      Claim 1;
                                                                                                                                                                                                                                                                    Novel isolated nucleic acid molecule associated with cancerous state of prostate cells and correlating with presence of prostate cancer, useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-FEB-2001;
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                                                                                                                                                                                   Page 3813-3814; 11750pp; English.
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The invention relates to an isolated nucleic acid molecule (I) com a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of t specification or its complement. (I) is useful for:
(a) assessing whether a patient is afflicted with prostate cancer;
(b) monitoring the progression of prostate cancer in a patient;
(c) assessing the efficacy of a test compound to inhibit prostate

) comprising of the

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RESULT 17
ABV28017
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(d) assessing the eff
in a patient;
(e) selecting a compo
(f) assessing the pro
(g) determining wheth
(h) assessing the agg
The invention relates to an isolated nucleic acid molecule (I) comprising a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the specification or its complement. (I) is useful for:
(a) assessing whether a patient is afflicted with prostate cancer;
(b) monitoring the progression of prostate cancer in a patient;
(c) assessing the efficacy of a test compound to inhibit prostate cancer in a patient;
(d) assessing the efficacy of a therapy for inhibiting prostate cancer in a patient;
(e) selecting a composition for inhibiting prostate cancer in a patient;
(f) assessing the prostate cell carcinogenic potential of a compound;
(g) determining whether prostate cancer has metastasized in a patient;
                                                                                                                                                                                                                                                                                                                                                    17-FEB-2000;
16-MAR-2000;
25-MAY-2000;
09-JUN-2000;
18-JUL-2000;
                                                                                                                                                                                                     Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, usefor detecting presence of prostate cancer, stage of prostate cancer
                                                                                                                                                                                                                                                             WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; prostate pharmacogenomic
                                                                                                                                                                             Claim 1; Page 5783-5785;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  assessing the efficacy of a therapy for inhibiting prostate cancer
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Conservative (
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RESULT 18
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Best Local S
Matches 16
                                                                                                                                                                                                                                                  The invention relates to a nucleic acid molecule (I) encoding a Can 1 polypeptide. The Can 1 nucleic acid molecule is used to diagnose or trinfertility or premature ovarian failure or Sertoli Cell only syndrome in a mammal. The present sequence is that of a human Can 1 encoding BAC clone RP11-334G22 of the invention.
                                                                                                                                                                    Sequence 180557
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; Can 1; antiinfertility; gynaecological; infertility; premature ovarian failure; menopause; Sertoli Cell only syn BAC clone RP11-334G22; GenBank reference AC007250; ds.
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                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page -; 45pp; English.
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  124080

    e: The present sequence is not given in the printed specification
obtained through the GenBank reference AC007250.

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Pred. No.
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                                                                                                          DB 24;
7.1;
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                                                                                                                          Length 180557;
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RESULT 19 AAH62689

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RESULT 20
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention provides the primary nucleotide sequence of the WSBV ge (AAH62689), predicted transcript sequences (AAH62689), predicted transcript sequences (AAH62689-AAH62839) and encoded proteins (AAG84910-AAG85051) and oligonucleotide sequences (AAH62840-63160) suitable for use as primers or probes. The nucleic a molecules and proteins of the invention are useful for diagnosis and monitoring viral infection, in screens for antiviral agents and for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Primary nucleotide sequence of the shrimp white spot Bacilliform virus (WSBV), useful for producing viral polypeptides that can be used to screen for agents that are useful for treating WSBV infection -
                                                                                                                                                                                                                                                                                                                                                                             monitoring viral gene expression or activity during a treatment regimen. The nucleic acid molecules are also useful as antisense constructs to control viral gene expression in infected cells and tissues and to create transgenic viral resistant shrimp.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antiviral agent; gene expression; antisense construct; transgenic viral resistant shrimp; ds.
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                                     gene therapy; chromosome
                                                                              Human secreted protein 5'
                                                                                                             06-OCT-2000
                                                                                                                                        AAC20071;
                                                                                                                                                                   AAC20071 standard; cDNA; 123
                                                                                                                                                                                                                                                                                                                                                   Sequence 305107 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Figure 2; 626pp; English.
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             Homo sapiens
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                                                     5' EST;
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THIRD INST OCEANOGRAPHY STATE OCEANI C SINOGENOMAX CO LTD.
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                                                                                                          (first entry
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                                                    expressed
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                                                                                                                                                                                                                                                                                                          100.0%;
                                                                                                                                                                                                                                                                                                          80.0%; Score 16; 100.0%; Pred. No.
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                                     sequence tag;
mapping; ss.
                                                                                 EST,
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                                                                                 SEQ ID NO: 24146
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                                                    secreted protein; cDNA isolation;
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                                                                                                                                                                                                                                                                                                          DB 22;
6.9;
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                                                                                                                                                                                                                                                                                                                       Length 305107;
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RESULT 21
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                              Addison's disease; allergy; asthma; osteoporosis; autoimmune thyroiditis; crohn's disease; dermatitis; diabetes; Graves' disease; haemodialysis; glomerulonephritis; scleroderma; systemic lupus erythematosus; uveitis; systemic sclerosis; ulcerative colitis; infection; trauma; pick disease; cardiovascular disorder; congestive heart failure; angina; epilepsy; hypertensive heart disease; neurological disorder; Alzheimer's disease; parkinson's disease; amyotrophic lateral sclerosis; stroke; dementia; Huntington's disease; multiple sclerosis; neuromuscular disorder; metabolic disorder; endocrine disorder; toxic myopathy; mental disorder; schizophrenic disorder; developmental disorder; anaemia; epilepsy; schizophrenic disorder; developmental disorder; anaemia; epilepsy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             identified within the present sequence. The 5' ESTs were prepared from total human RNAs or polyA+ RNAs derived from 30 different tissues. EST sequences usually correspond mainly to the 3' untranslated region (UTR) of the mRNA because they are often obtained from oligo-dT primed cDNA libraries. Such ESTs are not well suited for isolating cDNA sequences derived from the 5' ends of mRNAs and even in those cases where longer CDNA sequences have been obtained, the full 5' UTR is rarely included. 5' ESTs are derived from mRNAs with intact 5' ends and can therefore be used to obtain full length cDNAs and genomic DNAs. 5' ESTs are also used in diagnostic, forensic, gene therapy and chromosome mapping procedures. They are used to obtain upstream regulatory sequences and to design
                                                                                                                                                                                                                                                                                                             Secreted protein; SCEP; human; cell proliferative disorder; cancer; keratosis; arteriosclerosis; atherosclerosis; cirrhosis; hepatitis; psoriasis; autoimmune disorder; inflammatory disorder; AIDS; arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABK99951
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 123 BP; 48 A; 15 C;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and fo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2000-500381/45.
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                                                                                                                                                                                                                                                                                                                                                                                                        DNA encoding human secreted protein SCEP-34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABK99951 standard; DNA; 762
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence is one of a large number of 5' ESTs derived f mRNAs encoding secreted proteins. No ORF has yet been conclusively identified within the present sequence. The 5' ESTs were prepared
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-FEB-2000; 2000EP-0200610
                                                                                                                                                                                                                                                                                           acquired immunodeficiency syndrome; adult respiratory distress syndrome;
                                            hypothyroidism; glaucoma; sensorineural
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Pred. No.
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40;
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WO200248337-A2 Homo sapiens

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                                                                                                                                                                                                                                                                                                                                                                                 condition associated altered expression of functional SECP in a patient:

condition associated altered expression of functional SECP in a patient:

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condition associated altered expression of functional SECP in a patient:

condition associated altered expression of SECP in a subject or in a biological sequence of (I), and (II) and modulators of (I) are useful for disease treatment and prevention of cell proliferative disorders (e.g. cancer, condition) are useful for diagnosis, creatment and prevention of cell proliferative disorders (e.g. cancer, condition) and modulators of (I) are useful for diagnosis, creatment and prevention of cell proliferative disorders (e.g. cancer, condition) autoimmune/inflammatory disorders (e.g. acquired immune/inflammatory disorders (e.g. acquired commune/inflammatory disorders (e.g. acquired immune/deficiency syndrome, AIDS), adult respiratory distress syndrome, addition's disease, altergies, asthma, osteoporosis, autoimmune (conditions) disease, arthritis, scleroderma, systemic lupus erythematosus, systemic sclerosis, uncerative collitis, haemodialysis, uveitis; viral, bacterial, fungal, parasitic, protozoal, helminthic infections and creammand, cardiovascular disease), neurological disorders (e.g. Alzheimer's and plack uncertainty disease, amyotrophic lateral sclerosis, epilepsy, disease, parkinson disease, amyotrophic lateral sclerosis, epilepsy, and control and co
                                                                                                     Query Match
Best Local
                                                                                   Matches
                                                                                                                                                                                                                                 stroke, Huntington's disease, multiple sclerosis, dementia, neuromuscular disorders, metabolic, endocrine and toxic myopathies, mental disorders, schizophrenic disorders, and developmental disorders (e.g. anammia, epilepsy, hypothyroidism, glaucoma, sensorineural hearing loss and cataract). (II) is useful for creating transgenic animals to model human disease and to detect and quantity gene expression in biopsied tissues in which expression of SECP is correlated with disease. This sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel human secreted proteins and polynucleotides for diagnosing, preventing or treating cell proliferative, autoimmune/inflammatory, cardiovascular, neurological and developmental disorders -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                proteins (I), SECP 1-54. (I) and the polynucleotide encoding it useful for screening a compound for effectiveness as an agonist antagonist of (I) or compound that alters expression of (II). ()
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02-FEB-2001;
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05-JAN-2001;
                                                                                                                                                                      Sequence 762 BP; 174
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-DEC-2000;
555
                                      ຫ
                                                                                   15;
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Y J, Thangavelu K, Xu
BA, Honchell CD, Gietz
, Walla NK, Lu Y, Eli
Y, Khan FA, Tran
                                                                                                     Similarity
                                                                                                                                                                                                                   Ø
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INCYTE GENOMICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Page 221; 234pp;
                                        AGGAGGAACAAGTCC
                                                                                                                                                                                                                   human secreted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; 2001US-260105P.
; 2001US-262932P.
; 2001US-263096P.
                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2001US-263090P.
2001US-265926P.
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2000US-257852P.
                                                                                                     75.0%;
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541
                                                                                                                                                                                                               protein (SCEP).
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Y, Elliott VW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    isolated polypeptide chosen from secreted
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                                                                                                     Score 15;
Pred. No.
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                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Yue H, Ding L, La
Lee S, Tang YT,
KJ, Baughn MR, Ga
ott VW, Lu DAM, Ha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Yue H,
                                                                                                     36;
                                                                                                                              24;
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                                                                                                                           Length
                                                                                   Indels
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RESULT 22
ABZ66827
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ABL29548
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                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                 Best
                                                                                                                                                                                                                                                                                                                                   The invention relates to identifying compounds (A) with herbicidal activity from their ability to reduce or block expression or activity of gene products of specific nucleic acid sequences (I) or amino acid sequences (II) encoded by (I). (A) and antagonists of proteins encoded by (I) are useful as herbicides and plant growth regulators.

Overexpression of (I) in plants is used to produce herbicide-resistant plants and (I) may also be mutated to identify altered sequences that encode proteins resistant to herbicides. The present sequence is that of
                                                                                                                                                                                                                                                                                                   Sequence 1691 BP; 483 A; 309
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24-MAR-2003
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                      Drosophila; developmental pharmaceutical; gene; ds.
                                                           Drosophila
                                                                                                               ABL29548;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 154-155; 224pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13-FEB-2002; 2002WO-EP01466.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Arabidopsis thaliana.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Arabidopsis thaliana; herbicidal; plant; growth regulator; transgenic; herbicide-resistant; gene; ds.
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Drosophila melanogaster
                                                                                                                                      ABL29548 standard; DNA; 17255 BP
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23-MAY-2001; 2001DE-1025537.
                                                                                      26-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Identifying herbicides and plant growth regulators, from ability inhibit specific genes, also use of these genes to prepare
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-AUG-2002
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                                                                                                                                                                                                                                                               Local
                                                                                                                                                                                                                                                                                                                          polynucleotide of the invention.
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15; Conserv
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                                                                                                                                                                                                    TTAGGAGGAACAAGT 844
                                                            melanogaster genomic polynucleotide SEQ ID NO 40117.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Blau A,
                                                                                                                                                                                                                                                   Conservative
                                                                                      (first entry)
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                                                                                                                                                                                                                                                               100.0%;
                                                                                                                                                                                                                                                                           75.0%;
                                  biology; cell signalling; insecticide
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Pred. No.
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34;
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RESULT 24
ABQ76896
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Best Local Similarity
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                                                                                                                         Cos
                                                                                                                                                                                GPCR; protease; Parkinson's disc
single nucleotide polymorphism;
                                                                                                                                                                                   G-protein coupled receptor; secretin receptor subfamily; human; SNP; GPCR; protease; Parkinson's disease; gene; chromosome X; single nucleotide polymorphism; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to an isolated nucleic acid detection reagent capable of detecting 1000 or more genes from Drosophila. The inventueseful in developmental biology and in elucidating cell signalling cell-cell interactions in higher eukaryotes for the development of insecticides, therapeutics and pharmaceutical drugs. The invention discloses genomic DNA sequences (ABLIG176-ABL30511), expressed DNA sequences (ABLIG176-ABL30511), expressed DNA sequences (ABLIG176-ABL30511).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New isolated nucleic acid genes from Drosophila and interactions -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-MAR-2000; 2000US-191637P.
11-JUL-2000; 2000US-0614150.
                                                                                                                                                                                                                                                                                                            ABQ76896 standard; DNA; 53226
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17255 BP; 5336 A; 3353 C; 3445 G; 5121 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; SEQ ID NO 40117; 21pp + Sequence Listing; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Venter JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-MAR-2001; 2001WO-US09231.
                                                           variation
                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                    Human G-protein coupled receptor DNA SEQ ID
                                                                                                                                                                                                                                                            13-MAR-2003
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                        variation
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          replace (1746,c)

*tag= b

/note= "SNP, single nucleotide
replace (1755,g)

/*tag= c
/*tag= c
                                                                      /product= "GPCR"
/note "this coding
13 introns"
/note= "SNP, single nucleotide polymorphism"
                                                                                                                                  ocation/Qualifiers
                                                                                                                       000..50651
                                                                                                                                                                                                                                                                                                                                                                                                                        75.0%; Score 15;
100.0%; Pred. No
tive 0; Mismato
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29;
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                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                  polymorphism"
                                                                                   interrupted by
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/note= "SNP, single r
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/number= 5
16128..17484
/*tag= r
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/note= "SNP,
15911..16127
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/*tag= m

/number= 4

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/note= "SNP, single replace (24869,c)
                                  /note= "SNP, single replace (24774,a)
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/note= "SNP,
6038..6170
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*tar-
                                                                        /note= "SNP, single
replace (24206,g)
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/note= "SNP, single
replace (11390,t)
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/number= 3
1171..8059
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replace (5760,a)
                                                                                                          note= "SNP, single
replace (22935,t)
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replace (22910,t)
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17648..32332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /*tag= g
/number= 2
                                                                                                                                                                                                                                                                                               note= "SNP,
7485..17647
                                                                                                                                                                                                                                     number=
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/*tag= ar
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40089..42455

/*tag= as
/number= 10
42456..42577

/*tag= at
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/number= 13
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/number= 9
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/product= 8

36949..38129

/*tag= ao

/number= 8

/number= 8
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replace (25768,a)
/*tag= ac
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note= "a single nucleotide polymorphism (SNP)
    result in a deletion at this position"
    replace (29781,g)
                                                                                         14423..44691
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "SNP, single nucleotide polymorphism"
replace (28470..28471,t)
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/note= "SNP, single nucleotide polymorphism"
/eplace (26697,t)
                                                                                                     /*tag= av
/note= "a single nucleotide polymorphism (SNP)
/note= "result in a deletion at this position"
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/note= "SNP, single nucleotide polymorphism"
/eplace (30182,a)
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note= "SNP, single nucleotide polymorphism"
eplace (31772,t)
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eplace (28359,c)
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ceplace (26697,c)
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*tag= am
number= 7
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note= "SNP, single nucleotide polymorphism"
eplace (31936,g)
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note@ "SNP, single nucleotide polymorphism"
2333..32467
'tag= al
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   RESULT 25
AAQ76831
ID AAQ76831 standard; DNA; 125 BP.
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  Claim 1; Page 228; 616pp; English
                                                                                  Gross J, H
Sibson DR,
                                                                                                                                                                                                                                                                                                                                                       AAQ76831;
                                                            WPI; 1994-035056/04.
                                                                                                                                         13-JUL-1992;
                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                          Brain; placenta; bone marrow; genetic analysis; gene mapping; detection; homology; human; adrenal tissue; ds.
                                                                                                                                                                                                                                                                                                                    25-MAR-2003
23-SEP-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             exon
                                                                                                                                                                  13-JUL-1993;
                                                                                                                                                                                         20-JAN-1994
                                                                                                                                                                                                                                                                                             Human genome fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Webster M, Beasley EM, Ketchum KA,
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                        New nucleic acid fragment encoding for genetic analysis and mapping
                                                                                                                     (MEDI-) MEDICAL RES COUNCIL.
                                                                                                                                                                                                              WO9401548-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (WEBS/) WEBSTER M.
(BEAS/) BEASLEY E M.
(KETC/) KETCHUM K A.
(DFRA/) DI FRANCESCO V.
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                                                                                 Hadfield KM,
t, Starkey M;
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(first entry)
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/*tag= az
/number= 13
replace (48839,t)
/*tag= ba
/note= "SNP, single nucleotide polymorphism"
50267..50651
/*tag= bb
/number= 14
replace (52265,g)
/*tag= bc
/note= "SNP, single nucleotide polymorphism"
                                                                                                                                           92GB-0014857
                                                                                                                                                                  93WO-GB01467
                                                                                              Howells D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 15;
Pred. No.
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                                    gene
                                                                                              Kelly M,
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                                    products -
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RESULT 26
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Best Local S
Matches 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; GDP-mannose 4,6-dehydratase; GM4,6D; gene; ss; inflammation cellular fucosylation; glycoconjugate fucosylation; transplant rejearthritis; asthma; sepsis; reperfusion injury; stroke; infection; complex carbohydrate; gene replacement therapy; immunosuppressive; antiinflammatory; antiarthritic; antibacterial; cerebroprotective;
                                                                                 The invention relates to a composition comprising a human GDP-mannose 4,6-dehydratase (GM4,6D) peptide. The peptide is useful for identifying GM4,6D inhibitors are useful for reducing inflammation in a mammalian subject and for treating or ameliorating diseases affected by the level of cellular fucosylation or diseases affected by the fucosylation of glycoconjugates. These diseases include arthritis,
                                                                                                                                                                                                                                                                                                                                                                          New composition comprising GDP-mannose 4,6-dehydratase (GM4,6D) peptide, for manufacturing complex carbohydrates, or as targets for screening GM4,6D antagonists for treating e.g. arthritis, or transplant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABX30187 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    the placenta or bone marrow comprise any of: (A) a sequence selected from (AAQ76401-Q77613), (B) an allelic variation of a sequence as described in (A), or (C) a sequence complementary to (A) or (D)
                              transplant infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22-NOV-1996;
03-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-JUN-2001; 2001US-0878574.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human GDP-mannose 4,6-dehydratase (GM4,6D) DNA #12244.
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                                                                                                                                                                                                                                                                                              Disclosure;
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(Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GEMY)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2003-066673/06.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENETICS INST INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AGGAGGAACAAGTC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sequences exhibit no more than 90% homology to a human
rejection, asthma, sepsis, reperfusion injury, stroke or The GM4,6D peptide or a polynucleotide encoding it is also manufacturing complex carbohydrates and as targets for
                                                                                                                                                                                                                                                                                        SEQ ID NO 12246; 6pp; English
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97US-0984246.
98US-0149674.
99US-0333177.
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Pred. No.
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AAX993/C
ID AAX993
AC AAX933
AC CAncer
XW Luman
XW CAncer
XW Lransc
XW Wiln's
XW WY/rsp
XW WY/rsp
XW WY/rsp
XW Leukem
XW Drosta
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Best Local :
                                                                                                                                                                                                                            Crkvenjakov R, Dickson...

Bscobedo J, Garcia PD, Garcia v, Sones LW, Kassam A, Kennedy GC, K
Lamson G, Leshkowitz D, Pot D, Ra
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23-DEC-1997;
03-APR-1998;
21-OCT-1998;
27-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cancer; human; colon; breast; lung; transmembrane receptor; ATPase; integral membrane protein; aspartyl protease; GATA family; wnt family; transcription factor; G-protein alpha subunit; protein phosphatase; phorbolester binding protein; diacyllycerol binding protein; trypsin; protein kinase; tyrosine phosphatase; developmental signalling protein; www/rsp5/wwp domain; therapy; forensic; genetic mapping; diagnostic; detection; treatment; cervical; melanoma; colorectal adenocarcinoma; wilm's tumour; retinoblastoma; sarcoma; myosarcoma; lung carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        polynucleotide is useful in developing an assay for defects in the enzyme, as well as in gene replacement therapy. Sequences ABX17944 and ABX17947 ABX33716 represent DNA molecules encoding human GM4,6D peptides of the invention.

Note: The sequence data for this patent did not form part of the printed specification but was obtained in electronic format directly from USPTO
                                                                                                                                                                                                                                                                                                                                                                                                                       (CHIR )
(HYSE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     leukemia;
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                                                                                                                                                          isolated human polynucleotides
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               lymphoma; dysplasia; hyperplasia; endometrium; adrenal; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 98US-0217471.

97US-0068755.

98US-0080664.

98US-0105234.

98US-0105877.
                                                                                                                                                                                                                                                             Dickson M, Drmanac R, Drmanac S;

urcia PD, Garcia V, Giese K, Innis MA;

sam A, Kennedy GC, Kita D, Labat I;

kowitz D, Pot D, Randazzo F, Reinhard

Sudduth-Klinger J, Williams LT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98WO-US27610
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       70.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 14;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    58 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    77 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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This invention by screening for

on describes novel isolated human polynucleotides obtained for differential expression in colon cancer, breast cance ser cell lines. The polynucleotides of the invention are

Claim

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Page 348;

591pp; English

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2020202020202020202020202020202020
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cc cellular activities (AAA), eukaryotic aspartyl proteases, GATA family of Ct transcription factors, G-protein alpha subunit, phorbolesters or Cd diacylglycerol binding proteins, protein kinase, protein phosphatase 2C, CC protein tyrosine phosphatase, trypsin, wnf family of developmental .

CC signalling proteins and wW/rsp5/WWP domain containing proteins. The cencoded polypeptides also have a functional domain selected from Ank repeat, basic region plus leucine zipper transcription factors, concommental containing proteins. The commodomain, EF-hand, SH3 domain, WD domain/G-beta repeats, zinc finger (CCHC class), and zinc-binding metalloprotease (C (C2H2 type), zinc finger (CCHC class), and zinc-binding metalloprotease (C domain. The polynucleotides encode polypeptides with similarity to known protein families and are predicted to have similar properties. The novel protein familes can be used to develop products for use as therapeutic applications. In particular, the product can be used for tree detection and management of cancers. They can be used for treating e.g. cervical cancers, melanomas, colorectal adenocaroinomas, Wilm's tumour, sarcomas,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 1.4
                                                                                                                                                                         17-FEB-2000;
16-MAR-2000;
25-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                            Human; prostate cancer; cytostatic; carcinogen; pharmacodyanamic marker; pharmacogenomic marker; gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABV06501
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 retinoblastoma, myosarcomas, lung carcinomas, leukemias, such as chronic myelogenous leukemia, promyelocytic leukemia, monocytic leukemia, and myelocytic leukemia, and lymphomas such as histiccytic lymphoma, anhydric hereditary ectodermal dysplasia, congenital alveolar dysplasia, epithelial dysplasia of the cervix, fibrous dysplasia of bone, and mammary dysplasia, hyperplasias, e.g. endometrial, adrenal, breast, prostate or thyroid hyperplasias or pseudoepitheliomatous hyperplasia of prostate or thyroid hyperplasias or pseudoepitheliomatous hyperplasia of
               WPI; 2001-662795/76
                                                Schlegel R,
                                                                                                                                                            09-JUN-2000;
                                                                                                                                                                                                                                                20-FEB-2001; 2001WO-US05171
                                                                                                                                                                                                                                                                                      23-AUG-2001.
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                                                                                                                                                                                                                                                                                                                                                            HOMO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABV06501 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       proteins, 7 transmembrane receptors, ATPases associated with various
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        represented in AAX98275-X99118 and encode polypeptides of protein families selected from 4 transmembrane segments integral membrane
                                                                                                                                         18-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                            sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      230
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                                                                                    MILLENNIUM PREDICTIVE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CCTTAGGAGGAACA 217
                                                                                                                                       2000US-189862P.
2000US-207454P.
2000US-211314P.
2000US-219007P.
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                                             Endege
                                                                                                                                                                                                               2000US-183319P.
                                                                                                                        2000US-255281P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                    expression marker cDNA 6492
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              344
                                                Monahan JE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 14; pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                63 G;
                                                                                    MEDICINE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.3e+02;
hes 0;
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RESULT 29
AACO3189
ID AACO33
AC AACO3
AC AACO3
AC Humar
XW Humar
XW Gene
OX Homo
XX Ep10:
XX Ep10:
XX Ep10:
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Best Local S
Matches 14
                                                                                                                                                                                     New nucleic acid that is a 5' expressed sequence obtaining cDNAs and genomic DNAs that correspond diagnostic, forensic, gene therapy and chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 344 BP; 113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of t specification or its complement. (I) is useful for:
(a) assessing whether a patient is afflicted with prostate cancer;
(b) monitoring the progression of prostate cancer in a patient;
(c) assessing the efficacy of a test compound to inhibit prostate
                                                                                                                                                                                                                                                                                                                                                                           Dumas Milne Edwards
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EP1033401-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human secreted protein 5' EST, SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAC03189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to an isolated nucleic acid molecule (I) comprising a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel isolated nucleic acid molecule associated with cancerous state of prostate cells and correlating with presence of prostate cancer, useful for detecting presence of prostate cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-FEB-2000; 2000EP-0200610
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    in a patient
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (d) assessing the efficacy of a therapy for inhibiting prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         selecting a composition for inhibiting prostate cancer in a patient; assessing the prostate cell carcinogenic potential of a compound; determining whether prostate cancer has metastasized in a patient; assessing the aggressiveness or indolence of prostate cancer in a
                                                                                                                                                                                                                                                                                                                         2000-500381/45.
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                                                                                                                                                                                                                                                                                                                                                                                                                               GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                 AAG03183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AGGAGGAACAAGTC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EST;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AGGAGGAACAAGTC 113
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                                                                                                                                      ij
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99US-0122487
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                                                                                                                                 71pp
                                                                                                                                                                                                                                                                                                                                                                              Duclert A,
                                                                                                                                      + CD-ROM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        82 G; 94 T; 3 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                   Giordano
                                                                                                                                                                                           and chromosome
                                                                                                                                    English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.3e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein; cDNA isolation;
                                                                                                                                                                                                                 tag (5' EST) for
to 5'ESTs and for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                        mapping procedures
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sequence. derived f

The present sequence is one of a large number of 5' ESTs derived from mRNAs encoding secreted proteins. An ORF has been identified within the sequence. The 5' ESTs were prepared from total human RNAs or polyA+ RNAs

different tissues. EST sequences usually

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RESULT 30
AAS38136/c
ID AAS38136 standard; cDNA; 408
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Best Local S
Matches 14
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                 The invention relates to new polynucleotides and polypeptides, useful for diagnosis and treatment of breast, lung and colon cancer. The sequences can be used in detecting differentially expressed genes correlated with cancerous state of a mammalian cell, comprising detecting at least one differentially expressed gene product in a test sample derived from a cell suspected of being cancerous. They can also be used to inhibit tumour growth by modulating expression of a gene product. AAS36943-AAS39338 represent novel human diagnostic and therapeutic coding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Williams LT, Escobedo J, Innis MA, Garcia PD, Sudduth-Klinges
Reinhard C, Randazzo F, Kennedy GC, Pot D, Kassam A, Lamson
Drmanac R, Crkvenjakov R, Dickson M, Drmanac S, Labst I;
Leshkowitz D, Kita D, Garcia V, Jones WL, Stache-Crain B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 357
                                                                                                                                                                                                                                                                                  Claim 1; Page 896; 1193pp; English.
                                                                                                                                                                                                                                                                                                                                        New polynucleotides and polypeptides, useful for diagnosis treatment of breast, lung and colon cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-MAR-2000; 2000US-0188609
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel human diagnostic and therapeutic gene #1194.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17-DEC-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-530177/58
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (CHIR )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (HYSE-) HYSEQ INC.
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the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   breast;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      colon;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   prostate; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 21; 1
1.3e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      diagnostic;
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Query Match Best Local Similarity

Sequence 408

BP; 57

A; 110 C; 140 G; 101 T; 0 other;

70.0%; Score 14; DB 22; 100.0%; Pred. No. 1.3e+02;

Length 408;

07-NOV-2001 AAS26323;

(first entry)

AAS26323 standard; cDNA;

442

ΒP

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RESULT 31
AAX55494/c
ID AAX554
XX AAX554
XX AAX554
XX Pad1;
XX Pad1en
PI Polynu
XX (DUPO
XX Plant
PI Polynu
XX Plaim
XX Claim
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XX Claim
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XX Claim
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RESULT 32
AAS26323
ID AAS26
XX
AC AAS26
XX
DT 07-NC
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                                                                                                                                                                                                                                                                                                                                                     Best Local
                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention describes novel plant Padl, Crml or Jabl proteins which are capable of AP-1 transcription factor regulation. The proteins are thought to interact with transcription factors altering gene expression. The nucleic acid sequences of the invention may be used to isolate cDNAs and genes encoding homologous proteins from the same or other plant species. Synthetic peptides of the proteins may be synthesized to generate antibodies that are useful for screening expression libraries. Transgenic plants may be produced using the nucleic acid sequences to alter the levels of Padl, Crml and Jabl present in the plants. Altering the levels of these proteins would alter the level of AP-1 transcription factor activity in the plants.
                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 409 BP; 119 A; 90 C; 82 G; 118 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 7; Page 42; 57pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Plant homologues of yeast Pad1, polynucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Morgante M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-NOV-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-JUL-1999
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; soybean; wheat; rice; yeast; human; isolation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           91
                                                                                                                                                                                                                                                                                                                           14;
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                                                                                                                                                                                                                          CTTAGGAGGAACAA 283
                                                                                                                                                                                                                                                                         CTTAGGAGGAACAA 15
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       E I
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Human cDNA encoding Human; immunosuppre: cytostatic; cardian neuroprotective; an neuroprotective; an vulnerary; secreted hyperproliferative; cerebrovascular dis nervous system diso corneal infection; 2000US 24-FEB-2000; 2000US 24-FEB-2000; 2000US 21-JAN-2000; 2000US 21-MAR-2000; 2000US 21-MAR-2000; 2000US 21-MAR-2000; 2000US 21-MAR-2000; 2000US 21-JUL-2000; 2000US 21-JUL-2000; 2000US 21-JUL-2000; 2000US 21-JUL-2000; 2000US 21-AUG-2000; 2000US
human cDNA encoding a novel secreted protein, Seq ID 502.  human; immunosuppressive; antiarthritis; 8s; antitheumatic; varytosatalic; carddant; vasotropic; cerebropotectide; potential, vivulnerary; secreted protein; thematoid arthritis; wulnerary; secreted protein; thematoid arthritis; wulnerary; secreted protein; thematoid arthritis; propriotiserative disorder; cardiavascular disorder; cardia
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14-SEP-2000 14-SEP-2000 14-SEP-2000 14-SEP-2000 14-SEP-2000 14-SEP-2000 21-SEP-2000 21-SEP-2000 25-SEP-2000 25-SEP-2000 26-SEP-2000 29-SEP-2000 29-SEP-2000 29-SEP-2000 29-SEP-2000 29-SEP-2000 29-SEP-2000 20-OCT-2000 02-OCT-2000 03-NOV-2000 01-NOV-2000 01-DEC-2000 01-DEC-2000
2000US-0232397. 2000US-0232398. 2000US-0232401. 2000US-0233063. 2000US-0233064. 2000US-0233064. 2000US-0233064. 2000US-023364. 2000US-0234997. 2000US-0234997. 2000US-0234997. 2000US-0235336. 2000US-0235367. 2000US-0236367. 2000US-0236367. 2000US-0236367. 2000US-0236367. 2000US-0236367. 2000US-0236367. 2000US-0236370. 2000US-0236370. 2000US-0236370. 2000US-0236370. 2000US-0236370. 2000US-0236370. 2000US-0246476. 2000US-0246476. 2000US-0246476. 2000US-0246611. 2000US-0246278. 2000US-0246278. 2000US-0246278. 2000US-0246278. 2000US-0246278. 2000US-0246278. 2000US-0246278. 2000US-0249218. 2000US-0249219.

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                                                                                                                                                                                                                                                                                                                                                           CC rabbits, goats, horses, cates, dogs, chickens or sheep. They are also used in diagnosing a pathological condition or susceptibility ct to a pathological condition. Antibodies to the proteins can also compared to the proteins can also be used in alleviating symptoms associated with the disorders and in condition or susceptibility compared to the proteins can also be used in alleviating symptoms associated with the disorders and in compared to the proteins can also be used in alleviating symptoms associated with the disorders and in collide autoimmunosasays (EJISA). Disorders which are diagnosed or treated conclude autoimmune diseases e.g. rheumatoid arthritis, corebroular disorders e.g. cardiac arrest, cerebrovascular disorders correctly are cardiac arrest, cerebrovascular disorders correctly are cerebral ischaemia, angiogenesis, nervous system disorders e.g. corneal infection, and many other consideration occurs of the specification. The polypeptides can also be used to aid wound healing and epithelial cell proliferation, to correctly skin aging due to sunburn, to maintain organs before cransplantation, for supporting cell culture of primary tissues, to casa a food additive or preservative to increase or decrease storage casa a food additive or preservative to increase or decrease storage casa food additive or preservative to increase or decrease storage.
                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                     Query Match
Best Local S
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05-DEC-2000;
05-DEC-2000;
06-DEC-2000;
08-DEC-2000;
08-DEC-2000;
08-DEC-2000;
Human; gene; ds; neural disorder; immune system disorder; renal disorder; muscular disorder; respiratory disease; reproductive disorder; gastrointestinal disorder; pulmonary disorder; cardiovascular disorder; hyperproliferative disorder; inflammatory disease; allergic reaction;
                                                                                                                                                                                                                                                                                                                        capabilities, fat content, lipid, protein, carbohydrate, vitamins minerals, cofactors and other nutritional components. The present sequence encodes a novel secreted protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 encoded secreted proteins. The nucleic acids and proteins are used to prevent, treat or amediorate a medical condition in e.g. humans, mice, rabbits, goats, horses, cats, dogs, chickens or sheep. They
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New nucleic acid molecules encoding 461 human secreted proteins for diagnosing, preventing, treating or ameliorating medical conditions used as food additives or preservatives -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1; SEQ ID No 502; 980pp; English.
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P-PSDB; AAU16336.
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                                                                                               18-MAR-2003
                                                                                                                        ABX73664;
                                                                                                                                                 ABX73664 standard; DNA; 442 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         invention relates to isolated nucleic acid molecules and their
                                                                                                                                                                                                                    141
                                                                    novel
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                                                                                                                                                                                                                                                                                      Similarity
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2000US-025198.

2000US-0256719.

2000US-0251479.

2000US-0251856.

2000US-0251869.

2000US-0251869.

2000US-0251990.

2000US-0251990.

2000US-0251990.

2000US-0254997.
                                                                    polynucleotide #492.
                                                                                                                                                                                                                                                                         Conservative
                                                                                          (first entry)
                                                                                                                                                                                                                                                                                     70.0%;
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                                                                                                                                                                                                                                                                                     Score 14;
Pred. No.
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                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                      1.3e+02;
                                                                                                                                                                                                                                                                                                    DB 22;
                                                                                                                                                                                                                                                                        0,
                                                                                                                                                                                                                                                                                                 Length 442;
                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                        vitamins,
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                                                                                                                                                                                                                                                                        Gaps
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    New polypeptides and nucleic acids, useful in gene therapy for
                           P-PSDB; ABU55404.
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08-SEP-2000;
21-SEP-2000;
21-SEP-2000;
25-SEP-2000;
27-SEP-2000;
29-SEP-2000;
29-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-AUG-2000
01-SEP-2000
01-SEP-2000
01-SEP-2000
01-SEP-2000
01-SEP-2000
05-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-AUG-2000;
14-AUG-2000;
14-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 blood related disorder; cancer; immunosuppressive; antiinflammatory; cardiovascular; nephrotropic; cytostatic; antiallergic; thrombolytic; haemostatic; antiarteriosclerotic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   31-JAN-2000;
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WPI; 2003-147444/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US2002132753-A1
                           Rosen CA,
                                                                      (ROSE/)
                                                                                                                                                                                         20-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                         29-SEP-2000;
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                                                     ROSEN C A.
RUBEN S M.
BARASH S C
                           Ruben SM,
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2000US-229509P.
2000US-229513P.
2000US-231413P.
2000US-234223P.
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2000US-251856P.
2000US-251868P.
2000US-251869P.
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2000US-225267P.
2000US-225268P.
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2000US-236368P.
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2000US-236327P.
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2000US-234997P.
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2000US-225447P
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                             Barash
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RESULT 34
ABV36454
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CC sequences ABX73173-ABX74167 represent human novel polynucleotides of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local :
                                                                                                                                                                             17-FEB-2000;
16-MAR-2000;
25-MAY-2000;
09-JUN-2000;
18-JUL-2000;
13-DEC-2000;
                          Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, usefor detecting presence of prostate cancer, stage of prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                       Human; prostate pharmacogenomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       muscular, respiratory, reproductive, cardiovascular or renal disorders -
Claim 1; Page 7534; 11750pp; English
                                                                                                                    Schlegel R,
                                                                                                                                                                                                                                                                                      20-FEB-2001; 2001WO-US05171
                                                                                                                                                                                                                                                                                                                   23-AUG-2001.
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                                                                                                                                                                                                                                                                                                                                                                              Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human prostate
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABV36454 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 442
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                treating, inhibiting or preventing e.g. neural, immune system,
                                                                                                                                                  (MILL-)
                                                                                                                                                                                                                                                                                                                                                                             sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      141
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                                                                                                                                                  MILLENNIUM PREDICTIVE MEDICINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TAGGAGGAACAAGT 154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TAGGAGGAACAAGT
                                                                                                                                                                             ; 2000US-183319P.
; 2000US-189862P.
; 2000US-207454P.
; 2000US-211314P.
; 2000US-219007P.
; 2000US-255281P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                    expression marker cDNA 36445
                                                                                                                                                                                                                                                                                                                                                                                                     cancer; cytostatic; carcinogen; pharmacodyanamic marker;
marker; gene; ss.
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                                                                                                                       Monahan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gastrointestinal, pulmonary,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.3e+02;
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RESULT 35
ABA43976/c
ID ABA439
XX ABA439
XX ABA439
XX O1-FEB
XX Human
XX Human;
KW Human;
KW Giseas
XX Homo s
XX No-AUG
PR 21-SEE
PR 21-
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                              04-FEB-2000; 2000US-0180312.
26-MAY-2000; 2000US-0207456.
30-JUN-2000; 2000US-0608408.
03-AUG-2000; 2000US-0632366.
21-SEP-2000; 2000US-0234687.
27-SEP-2000; 2000US-0236359.
04-OCT-2000; 2000GB-0024263.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             patient;
(I) is a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to an isolated nucleic acid molecule (I) comprising a nucleotide sequence given in Tables 1-9 (ABV0010-ABV62213) of the specification or its complement. (I) is useful for:
(a) assessing whether a patient is afflicted with prostate cancer;
(b) monitoring the progression of prostate cancer in a patient;
(c) assessing the efficacy of a test compound to inhibit prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-FEB-2002
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                                                                                                                                                                                 New spatially-addressable set of single exon nucleic acid probes, useful for measuring gene expression in sample derived from human breast, comprises number of single exon nucleic acid probes
                                                                                                                                                                                                                                                                                          WPI; 2001-496933/54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human breast cell single exon nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cancer in a patient;
(d) assessing the efficacy of a therapy for inhibiting prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-JAN-2001; 2001WO-US00662
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200157271-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; microarray; single exon probe; gene expression; breast;
                                                                                                                                                                                                                                                                                                                                                                                                 (MOLE-) MOLECULAR DYNAMICS INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             is also useful as a pharmacodyanamic or pharmacogenomic marker
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14; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AGGAGGAACAAGTC 18
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                                                                                                                                   SEQ
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                                                                                                                                                                                                                                                                                                                                             Chen
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Pred. No.
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                                                                                                                                   sequence
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                                                                                                                                   listing; English
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The invention relates to a spatially-addressable set of single exon nucleic acid probes for measuring gene expression in a sample derived from human breast and BT 474 cells. The method involves contacting the probes with a collection of detectably labelled nucleic acids

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Minimum
Maximum
                                Result
No.
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Match
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18
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                                                                                             22781392 seqs,
94.4
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1080
124
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CB267362 1006268 H
BQ359040 PM3-GN051
BQ359216 PM3-GN051
BF850266 CM3-EN007
CB130696 K-EST0180
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RESULT 1
BF536318
LOCUS
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KEYWORDS
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Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Jeffrey E. Green, M.D.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
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602051968F1 NCI_CGAP_SG2
                                                                       Mammalia; Eutheria; Rodentia; Sciurognath
1 (bases 1 to 927)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                 mRNA sequence.
                                                                  Unpublished
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clone IMAGE: 4191040
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BG284498 602408640
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High quality sequence
Location/Qual
                                                                                                                                                                              Estimate of human gene number provided by using Tetraodon nigroviridis DNA sequence Nat. Genet. 25 (2), 235-238 (2000)
                                                                                                                                                                                                                                                                          AL188554.1 GI:7826658
GSS; genome survey sequence.
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/mol_type="mRNA"
  /organism="Tetraodon nigroviridis"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               strain="FVB/N"
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Tetraodon nigroviridis
Tetraodon nigroviridis
Tetraodon nigroviridis
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Neoteleostei;
Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;
Acanthomorpha; Acanthopterygii; Percomorpha; Tetraodontiformes;
Tetradontoidea; Tetraodontidae; Tetraodon.
Web: www.genoscope.cns.fr)
This sequence is a single read and was generated as scale clone-end sequencing project of the Tetraodon genome. For more information, please take a look at http://www.genoscope.cns.fr/Tetraodon.
                                                                                                                                                                          Direct Submission
Submitted (12-APR-2000) Genoscope - Centre National de Sequencage :
BP 191 91006 EVRY cedex - FRANCE (E-mail : seqref@genoscope.cns.fr
                                                                                                                                                                                                                                                                                                                                                                                    Roest Crollius,H., Jaillon,O., Dasilva,C., Ozouf-Costaz,C., Fizames,C., Fischer,C., Bouneau,L., Billault,A., Quetier,F., Saurin,W., Bernot,A. and Weissenbach,J. Characterization and repeat analysis of the compact genome of the freshwater pufferfish Tetraodon nigroviridis Genome Res. 10 (7), 939-949 (2000)
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Bernot, A., Fizames, C., Wincker, P., Brottier, P.,
Saurin, W. and Weissenbach, J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CNS02AHD 1080 bp DNA linear GSS 01-SEP-
Tetraodon nigroviridis genome survey sequence T7 end of clone
251P01 of library G from Tetraodon nigroviridis, genomic survey
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/lab_host="DH10B (T1 phage-resistant)"
/clone lib="NCI CGAP SG2"
/note="Organ: salivary gland; Vector: pCMV-SPORT6; Site_Not1; Site_2: Sal1; Cloned unidirectionally. Primer: Oli dT. Average insert size 1.3 kb. Constructed by Life Technologies. Note: this is a NCI CGAP Library."

224 c 266 g 214 t Z others
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r,P., Quetier,F.,
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                                                                                                                                                                                                                                                                                                                                                                          Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=MR4&t2=MR4-ET0142-
310101-001-h11&t3=2001-01-31&t4=1)
Seq primer: puc 18 forward
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Dias Neto, E., Garcia Correa, R., Verjovski-Almeida, S., Briones, M.R.,
Dias Neto, E., Garcia Correa, R., Verjovski-Almeida, S., Briones, M.R.,
Nagai, M.A., da Silva, W. Jr., Zago, M.A., Bordin, S., Costa, F.F.,
Goldman, G.H., Carvalho, A.F., Matsukuma, A., Baia, G.S., Simpson, D.H.,
Goldman, G.H., Carvalho, A.F., Matsukuma, A., Baia, G.S., Simpson, D.H.,
Brunstein, A., deOliveira, P.S., Bucher, P., Jongeneel, C.V., O'Hare
,M.J., Soares, F., Brentani, R.R., Reis, L.F., de Souza, S.J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar,
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tel: +55-11-2704922
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Simpson A.J.G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Shotgun sequenging of the human transcriptome with ORF expressed
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a 267 c 223 g 324 t
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               tissue mRNA and cDNA amplification low stringency conditions."
51 c 19 g 28 t
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Query Match

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RESULT 4
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                  Bource
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CE 1 (bases 1 to 170)

RS Adams, M.D., Kerlavage, A.R., Fleischmann, R.D., Fuldner, R.A., Bult (C.J., Lee, N.H., Kirkness, E.F., Weinstock, K.G., Gocayne, J.D., White, O., Sutton, G., Blake, J.A., Brandon, R.C., Man-Wai, C., Clayton, R.A., Cline, T.R., Cotton, M.D., Earle-Hughes, J., Fine, L.D., Fitzgerald, L.M., Fitzhugh, W.M., Fritchman, J.L., Geoghagen, N.S., Glodek, A., Gnehm, C.L., Hanna, M.C., Hedblom, E., Hinkle, P.S. Jr., Kelley, J.M., Kelley, J.C., Liu, L.-I., Marmaros, S.M., Merrick, J.M., Kelley, J.C., Liu, L.-I., Marmaros, S.M., Merrick, J.M., Shirley, R., Phillips, C.A., Ryder, S.E., Scott, J.L., Saudek, D.M., Shirley, R., Phillips, C.A., Ryder, S.E., Scott, J.L., Saudek, D.M., Shirley, R., Bedmarik, D.P., Cao, L., Cepeda, M.A., Coleman, T.A., Collins, E.J., Bedmarik, D.P., Cao, L., Cepeda, M.A., Fischer, C., Hastings, G.A., He, W.M., Hu, J.S., Greene, J.M., Gruber, J., Hudson, P., Kim, A.K., Kozak, D.L., Kunsch, C., Hungjun, J., Li, H., Meissner, P.S., Olsen, H., Raymond, L., Wei, Y.F., Wing, J., Xu, C., Yu, G.L., Ruben, S.M., Dillion, P.J., Fannon, W.R., Rosen, C.A., Haseltine, W.A., Fields, C., Fraser, C.M. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              90 GGCTGGTGTCACCTGT 75
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EST100142 Pancreas tumor
AA294899
1 GGCTGGTGTCACCTGT
                                                              l Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                     Email: arkerlav@tigr.org
For clone availability, additional sequence and exinformation related to this EST, please check the Index (http://www.tigr.org/tdb/hgi/hgi.html)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Other_ESTs: THC17274
Contact: Kerlavage,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The Institute for Genomic Research madical Center Drive, Rockville,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Initial assessment of human gene diversity and expression patterns based upon 83 million nucleotides of cDNA sequence Nature 377 (6547 Suppl), 3-174 (1995)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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EST.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bioinformatics
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                                                              88.9%;
ilarity 100.0%;
Conservative
                                                                                                                                                                      39
                                                                                                                                                                    /note="Organ: pancreas; Vector: pBluescript
ECORI; Site_2: XhoI"
48 c 51 g 28 t 4 others
                                                                                                                                                                                                                                                                                                                 /organism="Homo sapiens"
/mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                        /db_xref="taxon:9606"
/dev_stage="adult:
                                                                                                                                                                                                                                                                                                /db_xref="ATCC (inhost):190475"
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rlavage, AR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                additional sequence and expression his EST, please check the TIGR Human Gene
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Best Local (
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143
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FORWARD: 5'-GCCAAGCTGGAAATTAACCCTCACTAAAGGG-3'
BACKWARD: 5'-CCAGTGAATTGTAATACGACTCACTATAGGGCG-3'
Seq primer: 5'-GAAATTAACCCTCACTAAAGG-3'.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claudio, J.O., Masih-Khan, E., Tang, H., Goncalves, J., Voralia, M., Z.H., Nadeem, V., Cukerman, E., Franciso-Pabalan, O., Liew, C.C., Woodgett, J.R. and Stewart, A.K.

A molecular compendium of genes expressed in multiple myeloma alload 100 (6), 2175-2186 (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tel: (416) 946-4639
Fax: (416) 946-6546
Email: k.stewart@utoronto.ca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               610 University Ave., 5-126, Toronto, Ontario, M5G 2M9, Canada
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     University Health Network
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Oncology Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: A. Keith Stewart,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa;
Mammalia; Eutheria;
1 (bases 1 to 183)
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                                                                                                                   Similarity
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  GGCTGGTGTCACCTGT 128
                                            GGCTGGTGTCACCTGT
                                                                                           Conservative
                                                                                                                                                                                                              41
                                                                                                                                                                                                                                         /Gev_Stage="rlasma Cell leuxemia"
/clone lib="Myeloma (PCL) cDNA library"
/clone = TVector: Lambda Zap Express; Site 1: EcoRI; Site 2:
XhoI; mRNA was purified from plasma cell leukemia patient's peripheral blood containing >95% myeloma. An oligo d(T)18 primer containing XhoI restriction site was used to prime first strand synthesis using M-MLV reverse transcriptase. To protect the cDNAs from XhoI digestion in subsequent cloning step, the nucleotide analogue 5-methyl-dCTP was added to the nucleotide mixture and [a-32p]dATP was added to monitor the quantity and quality of first strand synthesis. After second-strand synthesis and blunting of cDNA termini, EcoRI adapters were ligated followed by kinase treatment and digestion with XhoI. The cDNAs were then size-fractionated using Sephacryl S-500 column and then ligated into EcoRI and Sephacryl S-500 column and then ligated into EcoRI and Sephacryl S-500 column and then ligated into EcoRI and brid digested Lambda Zap Express vector. The ligation product was packaged using Gigapack II packaging extract. The library had primary titre of approx. 1x106. Clones from the primary library were randomly selected for single pass
                                                                                                                                                                                                              sequencing."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /cell_type="myeloma"
/dev_stage="Plasma cell leukemia"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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                                                                                                                 100.0%;
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Pred. No.
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                                                                                           Mismatches
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О.
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52;
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                                                                                                                                    Length 183;
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KEYWORDS
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                                                               AUTHORS
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                                                                                                                                                                                                                                                                                                                                                              70 GGCTGGTGTCACCTGT 55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: cgapbs-r@mail.nih.gov
ECO RI site shown at the beginning of the sequence.
Tissue Procurement: Louis M. Staudt, M.D., Ph.D.
cDNA Library Preparation: M.B. Soares Lab
cDNA Library Arrayed by: M.B. Soares Lab
DNA Sequencing by: M.B. Soares Lab
Clone distribution: MGC clone distribution information
found through the I.M.A.G.B. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
1 (bases 1 to 221)
Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R.,
Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F.,
Goldman,G.H., Carvalho,A.F., Matsukuma,A., Bala,G.S., Simpson,D.H.,
Brunstein,A., deOliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare
                                                                                                                                                                                                                   BF354113
PM4-HT0726-270500-002-c07
BF354113
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1 (bases 1 to 199)

NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                          Homo sapiens
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                                                                                                 Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                   BF354113.1
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UI-HF-BL0-adc-h-06-0-UI.r1 NIH_I
                                                                                                                                                          Homo sapiens (human)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /cell_type="germinal center B cells"
/cell_line="MGC85"
/lab host="PH10B (LTI)"
/clone_lib="NIH MGC 37"
/clone="wector: pT773-Pac; Site_1: NotI; Site_2: Eco RI;
/note="wector: pT773-Pac; Site_1: NotI; Site_2: Eco RI;
/constructed from size fractionated cytoplasmic mRNA
(1.5-2.5kb). Directionally cloned. Cells provided by Louis
M. Staudt, Ph.D. Library preparation by Maria de Fatima
Bonaldo, Ph.D. and M. Bento Soares, Ph.D."

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/tissue_type="lymph"
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/db_xref="taxon:9606"
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Pred. No. 54;
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Homo sapiens cDNA clone
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1 (bases 1 to 273)

NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?t1=PM4&t2=PM4-HT0726-
270500-002-c07&t3=2000-05-27&t4=1)
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Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP,
                                                                                                        Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: CLONETECH Laboratories, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and Simpson,A.J.
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CDNA Library Preparation: CLONETECH Laboratories, Inc. CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                           Unpublished
                                                                                                                                                                                                                                                                                                                                 Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                    mRNA sequence.
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Location/Qualifiers
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/dev_stage="Adult"
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|mol_type="mRNA"
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RESULT 9
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                                                                                                                                          Other_ESTs: THC17274
Contact: Kerlavage,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Plate: LLCM815 row: k column:
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Adams, M.D., Kerlavage, A.R.,
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9712 Medical Center Drive, Rockville, Tel: 3018699056
                                                                           The Institute for Genomic Research
                                                                                                                  Bioinformatics
                                                                                                                                                                                                                                                                                                    based upon 83 million
Nature 377 (6547 Suppl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryola; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                            Initial assessment of human gene diversity and expression patterns
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
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/note="Organ: pancreas; Vector: pDNR-LIB (Clontecn);
/note="Organ: pancreas; Vector: pDNR-LIB (Clontecn);
/note="Organ: pancreas; Vector: policy of the Colonies o
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/clone="IMAGE:3949966"
/lab_host="DH10B (T1 phage-resistant)"
/clone_lib="NIH_MGC_78"
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                                                                                                                                                                                                                                                                                                    (6547 Suppl), 3-174 (1995)
                                                                                                                                                                                          THC172745
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58;
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186 GGCTGGTGTCACCTGT 171

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AU099544/c
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Best Local
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AUTHORS
                                                                                                                                                                                                                                                                     source
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         161 GGCTGGTGTCACCTGT 146
                                                                                                                                                                                                                                                                       Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and Sugano,S. Construction and characterization of a full length-enriched and a 5'-end-enriched CDNA library. Gene 200 (1-2), 149-156 (1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 (bases 1 to 300)
Suzuki,Y., Tsunoda,T., Taira,H., Mizushima-Sugano,J., Sese,J., Hata,H., Ota,T., Isogai,T., Tanaka,T., Nakamura,Y., Morishita,S., Okubo,K., Suyama,A. Pand.Sugano,S.
In silico mapping of the 5'-ends of human mRNAs using full-length enriched and 5'-end enriched cDNA libraries constructed by Oligo-capping method
                   1 GGCTGGTGTCACCTGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AU099544 300 bp mRNA linear EST 05-APR-200
AU099544 Sugano Homo sapiens cDNA library Homo sapiens cDNA clone
HSI08640 similar to Human translation initiation factor eIF3 p44
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       For clone availability, additional sequence and expression information related to this EST, please check the TIGR Human Gene Index (http://www.tigr.org/tdb/hgi/hgi/html)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: arkerlav@tigr.org
                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Yutaka Suzuki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens (human)
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sapiens
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                                                           Conservative
                                                                                                                                                        74
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /dev_stage="fetus, 24 wks"
/clone lib="retal brain I"
/note="Organ: brain; Vector:
EcoRI; Site 2: XhoI"
3 74 c 89 g 53 t
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                                                                                                                                                  /clone_lib="Sugano Homo sapiens cDNA library"
77 c 95 g 54 t
                                                                                                                                                                                        /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="HSI08640"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
/mal_type="mRNA"
/db_xref="ATCC (inhost):140673"
/db_xref="taxon:9606"
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                                                                           88.9%;
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                                                                           Score 16; DB 9; Length 300; Pred. No. 60;
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Pred. No.
                                                           Mismatches
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60;
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                                                         0; Indels
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AUTHORS
TITLE
 BASE COUNT
ORIGIN
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BF516504
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ACCESSION
VERSION
KEYWORDS
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DEFINITION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Robert Strausberg, Ph.D.

Email: cgapbs-r@mail.nih.gov
Oligo-dT track not found, Not I site shown in beginning of sequence
is likely internal to the message. cDNA Library Preparation: M.B.

Soares Lab Clone distribution: NCI-CGAP clone distribution
information can be found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BY515504 301 bp mRNA linear EST 07-DEC-:
UI-H-BW1-aoc-g-07-0-UI.81 NCI_CGAP_Sub7 Homo sapiens cDNA clone
IMAGE.3084469 3', mRNA sequence.
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Seq primer: M13 Forward
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National Cancer Institute, Cancer Genome Anat
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                                                                                                                        /Lab_nost="WHIUE (LIFE TECHNOLOGIES)
/clone lib="WCI CGAP Sub7"
/clone lib="WCI CGAP Sub7"
/clone lib="WCI CGAP Sub7"
is a subtracted library derived from NCI CGAP Sub6
NCI CGAP Sub7 library had 12 million recombinants. A
single-stranded DNA preparation of NCI CGAP Sub6 was used
as a tracer in a subtractive hybridization with a driver
comprising: the IMAGE pool (NCI CGAP Kid3 pool 1 LLAM
3334-3337, 3682-3683, 3798-3803 [IMAGE CloneIDs
1322376-1323911, 1456008-1456775,1500552-1502855);
NCI CGAP Kid5 pool 1 LLAM 3338-3342,3722-3725, 3776-3778
(IMAGE CloneIDs 1323912-1325831, 1471368-1472903,
1492104-1493255); NCI CGAP LUS pool 1 LLAM 3575-3582,
1851-3854 (IMAGE CloneIDs 1414920-1417991, 1520904-1522439); NCI CGAP EAS pool 1 LLAM 3682-3720,
3733-3735 (IMAGE CloneIDs 1257099-125861,1469064-1470983,
1475592-1476743); NCI CGAP PC2 pool 1 LLAM 2677-2459,
2758-2779, 362-368 (IMAGE CloneIDs 985608-986739,
1101192-1101959, 1217928-1220615); NCI CGAP CO10 pool 1
LLAM 2644-2653, 2871-2872 (IMAGE CloneIDs 1057416-1061255,
1144584-1143351); (6% of the driver population), plus a pool of 3,840 arrayed clones from NCI CGAP Sub3 (IMAGE CloneIDs 2708616-2712455)
CloneIDs 27036616-2712455) (4% of the driver population),
plus a pool of 5,472 clones from NCI CGAP Sub3 (IMAGE CloneIDs 271465-7712455)
plus a pool of 5,472 clones from NCI CGAP Sub3 (IMAGE CloneIDs 2723592-2729326) (40% of the driver population),
Subtraction was performed as previously described (Bonaldo, Lennon & Soares (1996): Normalization and Subtraction:
   44
TAG_SEQ=None found"
88 c 82 g
                                                                   Two Approaches To Facilitate Gene Discovery. Genome Research 6, 791-806.
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/clone="IMAGE:3084469"
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/mol_type="mRNA"
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CB267362/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FORWARD: CTCGGGAAGCGCGCCATTGTGTTGGT
BACKWARD: AATACGACTCACTATAGGGCGAATTGG
Seg primer: GTTGGTACCCGGGAATTC.
1 (bases 1 to 319)
Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R.,
Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F.,
Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tel: 410 706 1672
Fax: 410 706 1622
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 317)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           University of Maryland
660 Redwood St, HH497, Baltimore,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Gong Da-Wei
Division of Endocrinology, Diabetes and Nutrition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
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                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                        Homo sapiens
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llarity 100.0%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                             /tissue_type="Adipose"
/clone_lib="Human Fat Cell 5'-Stretch Plus
/note="Vector: lambdaTriplEx"
78 c 97 g 52 t 1 others
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   organism="Homo sapiens"

[mol_type="mRNA"]
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61;
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Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 319)
Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R.,
Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Costa,F.F.,
Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F.,
Solva,H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H.,
Brunstein,A., deOliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare
,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  High quality sequence start: 17
High quality sequence stop: 319.
Location/Qualifiers
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Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Brunstein, A., deOliveira, P.S., Bucher, P., Jongeneel, C.V., M.J., Soares, F., Brentani, R.R., Reis, L.F., de Souza, S.J.
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Fax: +55-11-2707001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BQ369218
Contact: Simpson A.J.G.
                                                                             sequence tag
Proc. Natl.
                                                                                                                             Shotgun sequencing of the human transcriptome with ORF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GGCTGGTGTCACCTGT 16
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79 c 92 g 54 t 1 others
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/clone_lib="GN0516"
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/db_xref="taxon:9606"
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Project. This entry can be seen in the following URL (http://www.ludwig.org.br/scripts/gethtml2.pl?tl=PM3&t2=PM3-GN0516-020701-031-g03&t3=2001-07-02&t4=1)
Seq primer: puc 18 forward
High quality semmence.
Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human
Project. This entry can be seen in the following URL
                                                                                                                             Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar,
                                                                                                                                                                                                                                                                                                                                       Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R., Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F., Goldman,G.H., Carvalho,A.F., Matsukuma,A., Bain,G.S., Simpson,D.H., Brunstein,A., daoliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 GGCTGGTGTCACCTGT 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16;
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1 (bases 1 to 335)
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BF850266
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Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar,
                                                                                        Tel: +55-11-2704922
                                                                                                                Brazil
                                                                                                                                                                                            Contact: Simpson A.J.G.
                                                                                                                                                                                                                                                                               sequence tags
                                                                                                                                                                                                                                                                                                Shotgun sequencing of the human transcriptome with ORF
                                                                                                                                                                                                                                                                                                                          Simpson, A.J.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens (human)
                                                                                                                                                                                                                                                             Proc. Natl. Acad. Sci. U.S.A. 97
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note="Organ: placenta_normal; Vector: puc18; Site_1: Smal; Site_2: Smal; A mini-library was made by cloning products derived from ORESTES PER (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under low stringency conditions."

79 c 92 g 54 t 1 others
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /dev_stage="Adult"
/clone_lib="GN0516"
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Pred. No.
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EN0077 Homo sapiens cDNA,
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JOURNAL
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AUTHORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNT
                                                                                                                                                                                                                                             source
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                                                                                                                                                                                                                                                                                                                                    52 Eoeun-dong Yuseong-gu,
Tel: +82-42-860-4470
Fax: +82-42-860-4409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (http://www.ludwig.org.br/scripts/gethtml2.pl?t1=CM3&t2=CM3-EN0077-
181100-489-d09&t3=2000-11-18&t4=1)
seq_primer: puc_18_forward
                                                                                                                                                                                                                                                                                                                                                                                    Genome Research Center
Korea Research Institute of Bioscience
                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Kim YS
                                                                                                                                                                                                                                                                                                                                                                                                                                                21C Frontier Korean 
Unpublished
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CB130696.1
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                                                                                                                                                                                                                                                                               High quality sequence stop:
                                                                                                                                                                                                                                                                                                Email: yongsung@mail.kribb.re.kr
Plate: 17 row: C column: 06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oh, K.J., Cheong, J.E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kim, N.S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens (human)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              quality sequence start: 52 quality sequence stop: 335. Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="Organ: lung_normal; Vector: pucl8; Site_1: SmaI; Site_2: SmaI, A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under low stringency conditions."
/clome_lib="LilsNU354"
/note="Organ: Liver; Vector: pCNS-D2; Site_l: EcoRI;
/note="Organ: Liver; Vector: pCNS-D2; Site_l: EcoRI;
Site_2: NotI; The poly (A)+ RNA was dephosphorylated with
bacterial alkaline phosphatase (BAP) and then decapped
                                                                                         /cell_type="Polygonal"
/cell_line="SNU-354"
                                                                                                                              tissue_type="Liver"
                                                                                                                                                                    /db_xref="taxon:9606"
/clone="L11SNU354-17-C06"
                                                                                                                                                                                                    organism="Homo sapiens"
|mol_type="mRNA"
                                                                                                                                                                                                                                                          ocation/Qualifiers
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/mol type="mRNA"
/db_xref="taxon:9606"
/dev_stage="Adult"
                                                                            Lab
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GI:28095037
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ne_lib="EN0077"
                                                                      host="Top10F'"
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E., Sohn, H.Y., Kim,
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Pred. No.
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Kim,J.M., Park,
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62;
                                                                                                                                                                                                                                                                                                                                                                         ience & Biotechnology
305-333, South Korea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          nRNA linear EST 29-JAN-2003 cDNA clone L11SNU354-17-C06 5',
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H69706/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hillier, L., Lennon, G., Becker, M., Bonaldo, M.F., Chiapelli, B., Chiasee, S., Dietrich, N., DuBuque, T., Favello, A., Gish, W., Hawkins, M., Hultman, M., Kucaba, T., Lacy, M., Le, M., Le, N., Mardis, E., Moore, B., Morris, M., Parsons, J., Prange, C., Rifkin, L., Rohlfing, T., Schellenberg, K., Soares, M.B., Tan, F., Thierry-Meg, J., Trevaskis, E., Underwood, K., Wohldmann, P., Waterston, R., Wilson, R. and Marra, M. Generation and analysis of 280,000 human expressed sequence tags Genome Res. 6 (9), 807-828 (1996)
                                                                                                                                                                                                                                                                                                                   High quality sequence stops: 345 Source: IMAGE Consortium, LLML free through LLML; contact the This clone is available royalty-free through LLML; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Insert Length: 821 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. Louis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         yr93a08.sl Soares fetal liver spleen INFLS Homo sapiens cDNA clone mags:212822 3', mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: est@watson.wustl.edu
Insert Size: 821
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Wilson RK
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                   quality sequence stop: 345.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     intact mRNA was ligated with DNA-RNA linker including ECORI site by treatment of T4 RNA ligase and the first strand cDNA was synthesized from oligo dT-selected mRNA by priming with dT-tailed vector. The dT-tailed vector was adjusted to have about 60nt. The cDNA vector was
                     /dev_stage="20 week-post conception fetus"
/lab_host="DH10B (ampicillin resistant)"
                                                                                             /db_xref="taxon:9606"
/clone="IMAGE:212822"
                                                                                                                                             db_xref="GDB:3785663"
                                                                                                                                                                        organism="Homo sapiens"

ol_type="mRNA"
                                                                          sex=#male"
                                                                                                                                                                                                                           .394
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lib="Soares
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SOURCE

COMMENT

Query Match

88.9%;

Score 16;

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BM789452/c
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Oh,K.J., Cheong,J.E., Sohn,H.Y., Kim,J.M., Park,H.S., Kim,S. and
Kim,Y.S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     K-EST0069040 S22SNU16 Homo sapiens cDNA clone S22SNU16-2-A08 5',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   52 Eoeun-dong Yuseong-gu,
Tel: +82-42-860-4470
Fax: +82-42-860-4409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Korea Research Institute of Bioscience & Biotechnology 52 Eoeun-dong Yuseong-gu, Daejeon 305-333, South Korea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Kim YS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BM789452.1 GI:19137684
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens (human)
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L6; Conservative
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                      111
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/clone_lib="S22SNU16"
/note="Organ: Stomach; Vector: pT7T3-Pac; Site_1: EcoRI;
/note="Organ: Stomach; Vector: pT7T3-Pac; Site_1: EcoRI;
Site_2: NotI; The S22SNU16 library was contributed by the
Soares laboratory and it was constructed as described by
Bonaldo, M.F., Lennon, G. and Soares, M.B. (1996), Genome
Research 6(9): 791-806. RNA was prepared from harvested
cells of SNU-16 culture. SNU-16 cell was obtained from
Korean Cell Line Bank (KCLB). SNU-16 was established from
accitic fluids of Korean patients by Park J.G. et al.
(1990), Cancer Res 50: 2773-2780."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              yongsung@mail.kribb.re.kr
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81 c 106 g 102 t 3 others
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'mol_type="mRNA"
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                                                                                                                                                                                                                                                          tissue_type="Ascites"
cell_type="Lymphoblast-like"
cell_line="SNU-16"
                                                                                                                                                                                                                                                                                                                                      one="S22SNU16-2-A08"
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          1009264 Human I
cDNA 5', mRNA :
CB270357
                                                                                   CB270357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
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602122957F1 NIH_MGC_56 Homo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        National Institutes of Health, Mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 (bases 1 to 404)
NIH-MGC http://mgc.nci.nih.gov/.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens (human)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can
                                                                                                                                                                                                                                                            GGCTGGTGTCACCTGT 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                           108
                                                                                                                                                                                                                                                                                                                                                                                                                      /tissue type="primitive neuroectoderm"
/lab host="primitive neuroectoderm"
/lab host="primitive neuroectoderm"
/clome_lib="NIH_MGC_56"
/clome_lib="NIH_MGC_56"
/note="Organ: brain; Vector: pDNR-LIB (Clontech); Site_1:
/note="Organ: brain; Vector: pD
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mol_type="mRNA"
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                               Fat Cell 5'-Stretch Plus cDNA Library Homo sapiens sequence.
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                                                                                                                                                                                                                                                                                                                                                          Length 404;
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                                                                                   EST 20-FEB-2003
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AUTHORS
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KEYWORDS
SOURCE
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AA135078/c
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Best Local Similarity
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                                                                                                                                                                                                                                                    Hillier, L., Lennon, G., Becker, M., Bonaldo, M.F., Chiapelli, B., Chiasee, S., Dietrich, M., Dubuque, T., Favello, A., Gish, W., Hawkins, M., Hultman, M., Kucaba, T., Lacy, M., Le, M., Mardis, E., Moore, B., Morris, M., Parsons, J., Prange, C., Rifkin, L., Rohlfing, T., Schellenberg, K., Soares, M.B., Tan, F., Thierry-Meg, J., Trevaskis, E., Underwood, K., Wohldmann, P., Waterston, R., Wilson, R. and Marra, M. Generation and analysis of 280,000 human expressed sequence tags Genome Res. 6 (9), 807-828 (1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 413)

Yang,R.-Z., Shuldiner,A. and Gong,D.-W.
EST analysis of human adipose gene expression
Unpublished
                                                                                                                          Contact: Wilson RK
Washington University Sch
4444 Forest Park Parkway,
Email: est@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                   Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              zo26d06.rl Stratagene
IMAGE:588011 5', mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conteat: Gong Da-Wei
Division of Endocrinology, Diabetes and Nutrition
University of Maryland
660 Redwood St. HH497, Baltimore, MD 21201, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens (human)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tel: 410 706 1672
Fax: 410 706 1622
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EST.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                (bases 1 to 428)
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/tissue_type="Adipose"
/clone_Tib="Human Fat Cell 5'-Stretch Plus
/note="Vector: lambdaTriplEx"
/note="Vector: lambdaTriplEx" 5 others
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/mol_type="mRNA"
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Pred. No.
                                                                                                                          School of Medicine way, Box 8501, St. !
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                428 bp mRNA linear colon (#937204) Homo sapiens sequence.
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65;
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                                                                                                                                  Louis,
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AW732816/c
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AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleosto Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 430)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC) Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AW732816 430 bp mRNA linear EST 21-APR-2000 bb14h10,y1 NIH MGC 21 Homo sapiens cDNA clone IMAGE:2962915 5' similar to TR:Ö14801 O14801 EUKARYOTIC TRANSLATION INITIATION FACTOR 3 SUBUNIT: [1] ;, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens (human)
                                                                                                                                                                                                                                                                                           High quality sequence stop: 412.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AW732816.1 GI:7633154
                                                                                                                                                                                                                                                                                                                                        found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.gov/image/html/iresources.shtml Seq primer: -40RP from Gibco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Insert Length: 1117
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                                                                                                                                                                                                                                                                                                                                                                                                           cDNA Library Preparation: Ling Hong/Rubin Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
CDNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: MGC clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GGCTGGTGTCACCTGT 52
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ilarity 100.0%;
Conservative
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Location/Qualifiers
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/lab_host="SOLR cells (kanamycin resistant)"
/clone_lib="Stratagene colon (#937204)"
/clone_Togan: colon; Vector: pBluescript SK-; Site_1:
/note="Togan: colon; Vector: pBluescript SK-; Frimer:
/note="Togan: colon; Vector: pBluescript SK-; Primer:
Cligo dT. T-84 colonic epithelial cell line. Average
oligo dT. T-84 colonic epithelial cell line. Average
insert size: 1.0 kb; Uni-ZAP XR Vector; ~5 adaptor
sequence: 5 GAATTCGGCACGAG 3' ~3' adaptor sequence: 5
CTCGAGTTTTTTTTTTTTTTTTTTTT 3'"

a 112 c 113 g 71 t 4 others
/tissue_type="choriocarcinoma"
/lab_host="PH10B (phage-resistant)"
/clone_lib="NIH MCC_21"
/note="Organ: placenta; Vector: pOTB7; Site_1: XhoI;
/site_2: EcoR1; cDNA made by oligo-dT prining:
Directionally cloned into EcoR1/XhoI sites using the
following 5' adaptor: GGCACGAG(G). Size-selected >50
                                                                                                                                                                     /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:2962915"
                                                                                                                                                                                                                                                                       1. .430
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/clone="IMAGE:588011"
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Pred. No.
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     Size-selected >500bp
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RESULT 23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 (bases 1 to 431)
Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R.,
Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F.,
Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H.,
Brunstein,A., deOliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare
,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BF850584.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PM1-EN0065-191100-007-h05 EN0065 Homo sapiens cDNA, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     (http://www.ludwig.org.br/scripts/gethtml2.pl?t1=PM1&t2=PM1-EN0065-
191100-007-h05&t3=2000-11-19&t4=1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tel: +55-11-2704922
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequence tags
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Shotgun sequencing of the human transcriptome with ORF expressed
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens (human)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GGCTGGTGTCACCTGT 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          119 a
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                        65
                   /note="Organ: lung_normal; Vector: puc18; Site_1: Smal; Site_2: Smal; A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 195,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under low stringency conditions."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies)."

112 c 125 g 74 t
                                                                                                                                                                                                              /db_xref="taxon:9606"
/dev_stage="Adult"
/clone_lib="EN0065"
                                                                                                                                                                                                                                                                                 /organism="Homo sapiens"
/mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                     .431
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Query Match Best Local Similarity

88.9%;

Score 16; Pred. No.

DB 10; 66;

Length 431;

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RESULT 25
CB145604/c
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TITLE
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                                                                                                                             AUTHORS
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CB145604
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1 (bases 1 to 431)

Yang,R.-Z., Shuldiner,A. and Gong,D.-W.
Territallysis of human adipose gene expression
Kim, Y.S.
21C Frontier Korean EST Project 2001
Unpublished
                                                                                 Homo sapiens

Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Bukaryota; Metazoa; Chordata; Craniata; Hominidae; Homo.

Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 433)

Kim,N.S., Hahn,Y., Oh,J.H., Lee,J.Y., Ahn,H.Y., Chu,M.Y., Kim,M.R.,

Kim,N.S., Cheong,J.E., Sohn,H.Y., Kim,J.M., Park,H.S., Kim,S. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 GGCTGGTGTCACCTGT 16
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Division of Endocrinology, Diabetes and Nutrition
University of Maryland
660 Redwood St, HH497, Baltimore, MD 21201, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                K-EST0200441 L11SNU35481 Homo sapiens cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FORWARD: CTCGGGAAGCGCGCCATTGTGTTGGT
BACKWARD: AATACGACTCACTATAGGGCGAATTGG
Seq primer: GTTGGTACCCGGGAATTC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: dgong@medicine.umaryland.edu
PCR PRimers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tel: 410 706 1672
Fax: 410 706 1622
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CB270361
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/mol_type="mRNA"
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100.0%; F1
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Pred. No. 66;
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VERSION
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                                                                                                                                                                                                                                                                                                                                                                                                       187
                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 GGCTGGTGTCACCTGT 16
                               Homo sapiens (human)
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Location/Qualifiers
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CB129626.1 GI:28093354
                                                                                                                                                                                                                 K-EST0179303 L11SNU354 Homo sapiens
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Plate: 13 row: C column: 11
      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Genome Research Center
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Site 2: NotI; The poly (A)+ RNA was dephosphorylated with
bacterial alkaline phosphatase (BAP) and then decapped
with tabacco acid pyrophosphatase (TAP). The decapped
intact mRNA was ligated with DNA-RNA linker including
EcoRI site by treatment of T4 RNA ligase and the first
strand cDNA was synthesized from oligo dT-selected mRNA by
priming with dT-tailed vector. The dT-tailed vector was
adjusted to have about 60nt. The cDNA vector was
circularized with E. coli DNA ligase after digestion of
EcoRI which site is also included in vector. An RNA strand
converted to a DNA strand by Okayama-Berg method. The
obtained cDNA vectors were used for transformation of
competent cells E. coli ToplOF by electroporation method.
The cDNA libraries constructed by this method are
full-length enriched cDNA library. After analyzing and
sequencing about 2,000 - 3,000 colonies in original cDNA
library, the abundant cDNAs were selected and amplified by
PCR reaction using vector region primer including T7
promotor as 5' primer and N(dT)14 as 3' primer. The PCR
products were used as template for synthesis of
biotinylated single stranded RNA by in vitro transcription
biotinylated single stranded RNA by in vitro transcription
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/lab host="Top10F'"
/clone_lib="LilSNU35481"
/clone_lib="LilSNU35481"
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100.0%; Pi
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_line="SNU-354"
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L11SNU354-16-B02 5',
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TITLE
                                                             AUTHORS
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Kim,N.S., Hahn,Y., Oh,J.H., Lee,J.Y., Ahn,H.Y., Chu,M.Y., Kim,M.R.,
Oh,K.J., Cheong,J.E., Sohn,H.Y., Kim,J.M., Park,H.S., Kim,S. and
Kim,Y.S.
21C Frontier Korean EST Project 2001
                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 436)

Kim,N.S., Hahn,Y., Oh,J.H., Lee,J.Y., Ahn,H.Y., Chu,M.Y., Kim,M.R., Oh,K.J., Cheong,J.E., Sohn,H.Y., Kim,J.M., Park,H.S., Kim,S. and Kim,Y.S.
                                                                                                                                                                                                                            K-EST0195953 L11SNU35481 Homo sapiens cDNA clone L11SNU35481-19-E11 CB142213
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unpublished
Contact: Kim YS
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Plate: 16 row: B column: 02
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Frontier Korean EST Project 2001
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/note="Organ: Liver; Vector: pCNS-D2; Site_1: EcoRI;
/note="Organ: Liver; Vector: pCNS-D2; Site_1: EcoRI;
Site_2: NotI; The poly (A)+ RNA was dephosphorylated wi
bacterial alkaline phosphatase (BAP) and then decapped
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cell_line="SNU-354"
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|e_lib="L11SNU354"
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Tel: +82-42-860-4470
Fax: +82-42-860-4409
                             Homo sapiens (human)
                                                                                          EST.
                                                                                                                    BF858511 438 bp mRNA linear RC1-FT0190-221100-021-c02 FT0190 Homo sapiens cDNA, BF858511.1 GI:12246255
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Plate: 19 row: E column: 11
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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Site 2: NotI; The poly (A)+ RNA was dephosphory.lated with
bacterial alkaline phosphatase (BAP) and then decapped
with tabacco acid pyrophosphatase (TAP). The decapped
intact mRNA was ligated with DNA-RNA linker including
EcoRI site by treatment of T4 RNA ligate and the first
strand cDNA was synthesized from oligo dT-selected mRNA by
priming with dT-tailed vector. The dT-tailed vector was
adjusted to have about 60nt. The cDNA vector was
adjusted to have about 60nt. The cDNA vector was
circularized with E. coli DNA ligase after digestion of
EcoRI which site is also included in vector. An RNA strand
converted to a DNA strand by Okayama-Berg method. The
obtained cDNA vectors were used for transformation of
competent cells E. coli ToplOF by electroporation method.
The cDNA libraries constructed by this method are
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PCR reaction using vector region primer including T7
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products were used as template for synthesis of
biotinylated single stranded RNA by in vitro transcription
broatting and stranded RNA by in vitro transcription
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               reaction. The synthesized RNA probes were hybridized with antisense single stranded cDNAs prepared from original liberary and incubated with avidin-gel. After removing DNA-RNA hybrids by centrifuge, the subtracted cDNA libraries were constructed by transformation of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           electroporation method.
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/lab_host="Top10F'"
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_xref="taxon:9606"
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100.0%;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 436;
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                                                                                                                                                                                     mRNA sequence.
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SOURCE
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1 (bases 1 to 438)

Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R.,

Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F.,

Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H.,

Brunstein,A., deOliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare
,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
                                                                                                                                                                                                                                                                                mRNA seq
BG104409
                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 447)

NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1. GGCTGGTGTCACCTGT 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?t1=RC1&t2=RC1-FT0190-
221100-021-c02&t3=2000-11-22&t4=1)
Seq primer: puc 18 forward
Use primer:
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
                                                                                                                                                                                                                                                                                                  Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
                                          Contact: Robert Strausberg, Ph.D.
                                                                Unpublished
                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                          BG104409.1 GI:12598251
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sequence tags
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                                                                                                                                                                                                              Homo sapiens (human)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /clone_lib="FT0190"
/note="Organ: prostate_tumor; Vector: puc18; Site_1: Smal;
/note="Organ: prostate_tumor; Vector: puc18; Site_1: Smal;
Site_2: Smal; A mini-library was made by cloning products
derived from ORESTES PCR (U.S. Letters Patent application
No. 196,716 - Ludwig Institute for Cancer Research)
profiles into the pUC 18 vector. Reverse transcription of
tissue mRNA and cDNA amplification were performed under
low stringency conditions."
a 110 c 133 g 80 t
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_xref="taxon:9606"
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ne_lib="FT0190"
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Pred. No.
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66;
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TITLE
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Best Local :
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                                                                                                                                                                                                                                                                                                                        Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
cDNA Library Preparation: CLONETECH Laboratories, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLA
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information
                                                                                                                                                                                                      Plate: LLCM868 row: j column:
High quality sequence stop: 449.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   601818761F1 NIH_MGC_58 Homo sapiens cDNA clone IMAGE:4044398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mRNA sequence.
BF130453
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                                                                                                                                                                                                                                                                             found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert Strausberg, Ph.D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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1 (bases 1 to 449)
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/lab_host="DH108 (phage-resistant)"
/clone_lib="NIH_MGC_20"
/clone_lib="NIH_MGC_20"
/note="Forgan: skin; Vector: pOTB7; Site_1: XhoI; Site_2:
/note="Forgan: skin; Vector: pOTB7; Site_1: XhoI; Site_2:
cloned into EcoRI/XhoI sites using the following 5'
adaptor: GGCACGAG(G). Size-selected >500bp for average
insert size 1.8kb. Library constructed by Ling Hong in
the laboratory of Gerald M. Rubin (University of
California, Berkeley) using ZAP-cDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies)."
29 a 116 c 134 g 68 t
                   /db_xref="taxon:9606"
/clone="IVAGE:4004398"
/tissue_type="hypernephroma"
/lab host="DH10B (T1 phage-resistant)"
/clone_lib="NIH_MGC_58"
                                                                                                                                  /organism="Homo sapiens"
/mol_type="mRNA"
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Vector: pDNR-LIB (Clontech); Site_1:
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AUTHORS
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Best Local Similarity
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601501730F1 NIH_MGC_70 Homo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
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National Institutes of Health, Mammalian Gene Collection
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1 (bases 1 to 452)
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Plate: LLAM9707 row:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                     GGCTGGTGTCACCTGT 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                           guality sequence stop: 452.
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Double-stranded cDNA was prepared from cell line RNA. 5'
and 3' adaptors were used in cloning as follows: 5'
adaptor sequence: 5'-CACGGCCATTATGGCC-3' and 3' adaptor
sequence: 5'-ATTCTAGAGGCCGAGGCGCGACATG-dT(30)BN-3'
(where B = A, C, or G and N = A, C, G, or T). Average
insert size 1.35 kb (range 0.9-4.0 kb). 15/15 colonies
contained inserts by PCR. This library was enriched for
full-length clones and was constructed by Clontech
Laboratories (Palo Alto, CA)."
                                                                                                                                                                                /note=Torgan: pancreas; Vector: pCMV-SPORT6; Site 1: Not1; Site 2: Sall; Cloned unidirectionally. Primer: Oligo dT. Average insert size 1.1 kb. Library constructed by Life Technologies."

117 c 133 g 75 t
                                                                                                                                                                                                                                                                      /db xref="taxon:9606"
/clome="IMAGE:3903510"
/tissue_type="epithelioid carcinoma"
/lab host="DH10B (phage-resistant)"
/clome_lib="NIH_MGC_70"
                                                                                                                                                                                                                                                                                                                                                              /organism="Homo sapiens"
/mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                      ocation/Qualifiers
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Pred. No. 67;
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Pred. No.
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one IMAGE:3903510
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VERSION KEYWORDS

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JOURNAL COMMENT
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AUTHORS
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BX475327/c
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VERSION
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Best Local S
Matches 16
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                                                                                                                                                                                                                                                                          AA496839
453 bp mRNA linear EST 12-AUG-1997
ae33b02.rl Gessler Wilms tumor Homo sapiens cDNA clone IMAGE:897579
5' similar to contains Alu repetitive element;contains element TAR1
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 453)

Hillier,L., Allen,M., Bowles,L., Dubuque,T., Geisel,G., Jost,S., Kucaba,T., Lacy,M., Le,N., Lennon,G., Marra,M., Martin,J., Moore,B., Schellenberg,K., Steptoe,M., Tan,F., Theising,B., White,Y., Wylie, T., Waterston,R. and Wilson,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This clone (DKFZp686109176) is available at the RZPD in Berlin. Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059 Berlin- Charlottenburg, GERMANY; Email: clone@rzpd.de. Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Clone from S. Wiemann, Molecular Genome Analysis, German Cancer Research Center (DKFZ); Email s.wiemann@dkfz- heidelberg.de; sequenced by Qiagen (Hilden/Germany) within the cDNA sequencing consortium of the German Genome Project.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ingolstaedter Landstr.1, D-85764 Neuherberg, This is the 5' sequence of the clone insert
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EST (Bahr,A., Lauber,J., Mewes,H.W., Weil,B.,
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DKFZp686I09176 rl 686 (synonym: h

DKFZb686I09176 rl 686 (synonym: h

DKFZb686I09176 rl 686 (synonym: h
                                                                                                                                                                Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa;
Mammalia; Eutheria;
1 (bases 1 to 452)
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/note="Vector: pTriplEx2; Site_1: SfiIA; Site_2: SfiIB;
cDNA-collection"
a 120 c 138 g 80 t
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/mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /dev_stage="adult"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone="DKFZp686109176"
/tissue_type="human skeletal muscle"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          db_xref="taxon:9606"
                                                                                                                                                                                                                                                              element ;, mRNA sequence.
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Lawrence Livermore National Laboratory
Livermore, CA, USA
Email: help@image.llnl.gov
This read has been verified (found to hit its original self in the correct orientation), as part of the I.M.A.G.E. Consortium quality control effort. High quality sequence is defined as having 100 or more base pairs with a phred quality value of 20 or greater, where a sliding window of 4 base pairs with a phred quality value of 15 or greater marks the beginning and end of the sequence. For information on obtaining this clone, please contact
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Fax: 327 200 200 Email: est@watson.wustl.edu
Email: est@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
This clone is available royalty-free through LLNL; contact the
This clone is available royalty-free through LLNL; contact the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       454 bp · mRNA linear EST 04-DEC-20 imageqc 3_2001/smm196bdff41.xl NIH_MGC_61 Homo sapiens cDNA clone IMAGE:4686857 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Prange CK
The I.M.A.G.E. *Consortium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished Other_ESTs: BG529561
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 454)

Kale, P.I., Harsch, T.J., Folta, P.A., Nelson, D.O., Sanders, C.G. and
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Fax: 314 286 1810
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             resequencing for verification
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was prepared from a pool of 6 anonymous wilms' tumor RNAs.
RNA was prepared by acid-phenol, followed by one round of
oligo dT selection. cDNA library preparation was with
the BRL/Life Tech. Superscript Plasmid system. An
oligo-dT NotI primer for first strand system. An
oligo-dT NotI primer for first strand systems. An
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                                                                                                                                            found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Plate: LLAM9593 row: l column: 07
                                                                                                                                                                                                                                           Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
                                                                                                                                                                                                                                                                                    Unpublished
                                                                                                                                                                                                                                                                                                   NIH-MGC http://mgc.nci.nih.gov/
National Institutes of Health,
                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                          Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                   DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
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/note="Organ: ovary; Vector: pCMV-SPORT6; Site_1: Not1; Site_2: Sall; Cloned unidirectionally. Primer: Oligo dT. Average insert size 1.8 kb. Library constructed by Life BASE COUNT 129 a 117 c 133 g 76 t

Ouery Match 88.9%; Score 16; DB 10; Length 455; Matches 16; Conservative 0; Mismatches 0; Indels 0; Gaps 0; Matches 16; Conservative 0; Mismatches 0; Indels 0; Gaps 0; 1 GGCTGGTGTCACCTGT 16

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Search completed: August 15, 2003, 10:57:39

Job time: 1136.8 secs. 3.
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                                                                                                                                                                                                                                                                                                                                                                                                                JOURNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AUTHORS
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The sequence data in this record represents an 'enhanced' version of a Phase 2 submission. Specifically, the indicat order and orientation of each sequence contig has been established using one or more of the following: read-pair data from individual subclones, overlaps with neighboring clones, alignment with available reference sequence (e.g., human), and/or confirmation by PCR testing. In addition, the sequence assembly is based on at least 8X average
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Benjamin, B., Blakesley, R. W., Bouffard, G.G., Brinkley, C., Brooks, S. Cariaga, K., Coleman, B., Engle, J., Granite, S., Guan, X., Gupta, J., Haghighi, P., Han, J., Hansen, N., Ho, S.-L., Idol, J.R., Karlins, E., Laric, P., Lee-Lin, S.-Q., Legaspi, R., Maduro, Q.L., Maduro, V.B., Margulies, E.H., Masiello, C., Maskeri, B., McDowell, J., Paguirigan, C., Pearson, R., Portnoy, M.E., Prasad, A., Reddix-Dugue, N., Schandler, K., Schueler, M.G., Staon, C., Stantripop, S., Thomas, J. W., Thomas, P.J., Touchman, J. W., Vogt, J.L. Wetherby, K.D., Wiggins, L., Young, A. and Green, E.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AC137657.2 GI:29135576
HTGS_PHASE2; HTGS_DRAFT
Sus scrofa (pig)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AC137657
AC137657.2
                                                                                                                                                                                                                                                                                                                                                Submitted (21-MAR-2003) NIH Intramural Sequencing Center, 8717 Grovemont Circle, Gaithersburg, MD 20877, USA On Mar 21, 2003 this sequence version replaced gi:25700398.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Direct Submission
Submitted (27-NOV-2002) NIH Intramural Sequencing
Grovemont Circle, Gaithersburg, MD 20877, USA
3 (bases 1 to 175005)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Quantitative epstein barr virus pcr rapid assay
Patent: WO 02064842-A 1 22-AUG-2002;
Children's Hospital Research Foundation (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AC137657 175005 Nus scrofa clone RP44-494H16,
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Green, E.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Cetartiodactyla; Suina; Suidae; 1 (bases 1 to 175005)
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Green, E.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (bases 1 to 175005)
                                                                                                                                                                                                                                                         Center: NIH Intramural Sequencing Center code: NISC Web site: http://www.nisc.nih.gov Contact: nisc_zoo@nhgri.nih.gov
                                                                                                                                                                                               Center project name: ecq
Center clone name: 494H16
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                                                                                                                                                                                                                                         ----- Project Information
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/mol_type="genomic DNA"
/db_xref="taxon:10376"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
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Pred. No. 0.81;
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9
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C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Vogt, J.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Brooks, S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           8717
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1 8074: contig of 8074 bp in length

8075 3811: contig of 25637 bp in length

8175 3811: contig of 25637 bp in length

33812 33911: gap of unknown length

33912 36676: contig of 2765 bp in length

36677 50073: contig of 13297 bp in length

50174 50173: gap of unknown length

50174 57164: contig of 6991 bp in length

57165 59245: contig of 2181 bp in length

59446 59545: gap of unknown length

59446 59545: gap of unknown length

59546 59773: contig of 2181 bp in length

59546 59773: contig of 1329 bp in length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            is believed to be correct as given, however the sizes of the gaps between them are based on estimates that have provided by the submittor.

This sequence will be replaced by the finished sequence as soon as it is available and by the finished sequence as soon as it is available.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            consists of 12 contigs. Gaps between the contigs are represented as runs of N. The order of the pieces
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NOTE: This is a 'working draft' sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       124523
124623
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167608
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143906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      igs have been trimmed away, and each base is associated a Phrap-derived quality score.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequencing vector: plasmid; n/a; 100% of reads Chemistry: Dye-terminator Big Dye; 100% of reads Assembly program: Phrap; version 0.990319
Consensus quality: 171640 bases at least Q40
Consensus quality: 172768 bases at least Q30
Consensus quality: 173465 bases at least Q20
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72774
72874
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Insert size: 173905; sum-of-contigs
Quality coverage: 9.32x in Q20 bases;
Quality coverage: 9.01x in Q20 bases;
                                                                                                                                                                                                                                                  33912.
                                                                                                                                                                                                                                                                                                               vector_
                                                                 /note="assembly_fragment"
59546. .72773
                                                                                                           /note="assembly_fragment" 57265. .59445
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                                                                                                                                                                                                      36777.
                                                                                                                                                                                                                                                                                                                                      clone_end:T7
                                                                                                                                                                                                                                                                                                                                                       notes "assembly_fragment
                                                                                                                                                                                                                                                                                                                                                                                                          clone_lib="RP44"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   organism="Sus scrofa"
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note="assembly_fragment"/
                                       note="assembly_fragment"
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                                                                                                                                                                                                                                                                                                                                                                                                                                 clone="RP44-494H16"
                                                                                                                                                                                                                                                                                                                                                                                                                                                      db_xref="taxon:9823"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mol_type="genomic DNA"
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    Summary Statistics

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175005: contig of 7298
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143805: contig
                                                                                                                                                                                                      .50073
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g of 5247 bp in 1
f unknown length
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sum-of-contigs
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                                                                                                                                                                            The following abbreviations are used to associate primary accession numbers given in the feature table with their source databases:

Bm:, RMBL; Sw:, SWISSPROT; Tr:, TREMBL; Wp:, WORMPEP; Information on the WORMPEP database can be found at http://www.sanger.ac.uk/Projects/C_elegans/wormpep This sequence was generated from part of bacterial clone contigs of human chromosome 6, constructed by the Sanger Centre Chromosome 6 Mapping Group. Further information can be found at http://www.sanger.ac.uk/HGP/Chr6
http://www.sanger.ac.uk/HGP/Chr6
http://www.sanger.ac.uk/HGP/Chr6
park Cancer Institute by the group of Pieter de Jong. For further details see http://bacpac.med.buffalo.edu/
VECTOR: pCYPAC2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Submitted (31-OCT-2000) Sanger Centre, Hinxton, Cambridgeshire, CB10 18A, UK. E-mail enquiries: humquery@Banger.ac.uk Clone requests: clonerequest@Banger.ac.uk clone requests: clonerequest@Banger.ac.uk clone requests: clonerequest@Banger.ac.uk clone on the sequence assembly data is compared from overlapping clones. During sequence assembly data is compared from overlapping clones. Where differences are found these are annotated as variations together with a note of the overlapping clone name. Note that the variation annotation may not be found in the sequence submission corresponding to the overlapping clone, as we submit sequences with a small overlap as described above.
                IMPORTANT: This sequence is not the entire insert of clone RP4-579A14 It may be shorter because we sequence overlapping sections only once, except for a 100 base overlap. The true left end of clone RP4-579A14 is at 1 in this sequence true left end of clone RP1-230L10 is at 88458 in this sequence. true right end of clone RP5-826L7 is at 60956 in this sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This sequence has been finished according to sequence map criteria as follows. An attempt is made to resolve all sequencing problems, such as compressions and repeats, but not necessarily within known annotated human repeat sequence elements (e.g. Alu). Where the sequence is ambiguous, there is an annotation using the 'unsure'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
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167708. .175005
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12836. 13137
/note="Allusx repeat: matches 1...
13782. 14092
/note="Allus repeat: matches 1...3
14290. 14600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note="1318 bases of transposon Tn10 (J01829) removed here. This sequence represents the duplicated flanking sequence of the transposon."
7142..7177
                                                                                                                                                             note="AluSg repeat: matches 1.
15665. .15962
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/note="36 cc
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/note="14 co
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
/chromosome="6"
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                                                                                                                                          note="AluSx repeat: matches 1.
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note="L1PA13 repeat: matches 5663. .6149 of consensus"
0302. .10348

note="MER91A repeat: matches 120. .166 of consensus"
0349. .10724

note="MSTA repeat: matches 1. .426 of consensus"
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78. .1087
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     note="9 copies 4 mer acac 88% conserved"
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note="8 copies 19 mer 89% conserved"
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10te="HAL1 repeat: matches 700. .906 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    note="16 copies 2 mer at 87% conserved"
                  10te="MSTA repeat:
1286: 17448
                                                                                                                                                                                                                                                                                                                                            ote="MER57B repeat:
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ote="MER5A repeat: matches 1. .189 of consensus"
48. .4357
ote="MIR repeat: matches 94. .204 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                te="MLT1A1 repeat: matches 1. .365 of consensus" 85. .11195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6. .8248
te="AluSx repeat: matches 40. .280 of consensus"
3. .8684
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          te="36 copies 17 mer 80% conserved"
2. .4061
ce="MIR repeat: matches 204. .243 o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              e="MSTB repeat: matches 1. .426 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                             e="MER91A repeat: matches 1. .114 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             e="MLT1C repeat: matches 22. .455 of consensus"
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                                                                 repeat: matches 176.
repeat: matches 32. .202 of consensus"
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                               matches 1. .130 of consensus"
                                                                                                                                                                                                                                                  matches 1. .311 of consensus"
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                                                                                                      matches
                                                                                                      5319. .5572
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                  /note="MLT1A1 repeat:
35465. .35756
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19387. .19574
/note="2 copies 94 mer 87% conserved"
                                                                                     note="AluY repeat: matches 7.
4594. .34840
                                                                 note="MLT1E repeat:
                                                                                                                                                                          note="MLT1H repeat: matches 112. .532 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            note="MLT1F repeat: matches 1. .228 of co
6108. .26267
note="40 copies 4 mer tgtg 65% conserved"
                                                                                                                                                                                                                                                                                      )089. .30632
note="match: GSS: Em:AQ318912"
                                                                                                                                          ote="AluY
                                                                                                                                                                                                                                                                                                                                                                            mplement (29578. .30078)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Note="MIR repeat: matches 40. .190 of consensus" 1272. .28443 tote="MIR repeat: matches 16. .208 of consensus" 1218. .29413
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    tote="59 copies 4 mer tgtg 71% conserved"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            109. .26268
Note="80 copies 2 mer gt 65% conserved"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           896. .24202
bote="MLT1F repeat: matches 228. .541 of consensus"
1203. .24563
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .445. .21527
lote="MLT1A1 repeat: matches 405. .501 of consensus"
                                                                                                                                                                                                                 ote="16
                                                                                                                                                                                                                                               ote="AluY repeat: matches 1. .306 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ote="45 copies 2 mer tt 71% conserved"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ote="6 copies 19 mer 68% conserved" 414. .29497
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ote="3 copies 94 mer 73% conserved"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 281. .26516
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ote="L2 repeat: matches 2425. .2664 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ote="AluSg_repeat: matches 1.
1796. .23027
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                                                                                                                                                                                                                                                                                                                        ote="MER58A repeat: matches 175. .224
                                                                                                                                                                                                                                                                                                                                                                                                                                  ote="5 copies 17 mer 80% conserved"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ote="21 copies 4 mer cttt 85% conserved"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   131. .29515
                                                                                                                                                                                                                                                                                                                                        Te="match: GSS: Em:AQ894713"
11. .29870
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e="match: GSS:
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                                                                                                                         .34569
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                                                                                                                                                                                                               copies 4 mer agag 82% conserved"
                                                                                                                                        repeat: matches 1. .305 of consensus"
repeat: matches 4.
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                                                                     matches 308.
                                 matches 65.
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AC128140/c
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Matches 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lotus japonicus
Lotus japonicus
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Spermatophyta; Magnoliophyta; Fabaceae; Papilionoideae; Loteae;
                                                                                                                                                                                                                                      l Similarity
17; Conserv
AC128140
AC128140.2 GI:23265194
HTG; HTGS_PHASE1; HTGS_DRAFT; HTGS_ENRICHED.
Rattus norvegicus (Norway rat)
                                                                                                                                                                                                         N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (12-DEC-2002) Shusei Sato, Kazusa DNA Research I. Department of Plant Gene Research; 2-6-7 Kazusa-kamatari, Chiba 292-0818, Japan (E-mail:ssato@kazusa.or.jp, URL:http://www.kazusa.or.jp/, Tel:81-438-52-3935(ex.2337),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kaneko, T., Asamizu, E., Kato, T., Sato, S., Nakamura, Y. and Tabata, S. Structural Analysis of a Lotus japonicus Genome. III. Sequence Features and Mapping of Sixty-two TAC Clones Which Cover the 6.7 M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  107979
Lotus japonicus genomic DNA,
complete servere
                                                                AC128140 226979 bp D
Rattus norvegiçus clone CH230-63023,
***, 2 unordered pieces.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Direct Submission
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DNA Res. 10, 27-33 (2003)
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                                                                                                                                                                                                                                                                                                                                                                             /organism="Lotus japonicus"
/mol_type="genomic DNA"
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/chromosome="3"
                                                                                                                                                                                                                                                                                                                /clone="LjT34M22"
/clone_lib="LjT library"
/note="TAC clone:TM0282"
21293 c 19709 g 33736 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        location/Qualifiers
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chromosome
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Anyalebechi, V., Aoyagi, A., Ayodeji, M., Baca, E., Baden, H.,
Baldwin, D., Bandaranaike, D., Barber, M., Barnstead, M., Benn
Biswalo, K., Blair, J., Blankenburg, K., Blyth, P., Brown, M.,
Bryant, N., Buhay, C., Burch, P., Burrell, K., Calderon, E.,
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                         Submitted (21-SEP-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
On Sep 21, 2002 this sequence version replaced gi:21908739.
The sequence in this assembly is a combination of BAC based reads and whole genome shotgun sequening reads assembled using Atlas (http://www.hgsc.bgm.tmc.edu/projects/rat/). As a result, the
                                                                                                                                                                                                                                                                                                                                                   Submitted (19-JUL-2002) Human of Molecular and Human Genetic Baylor Plaza, Houston, TX 7703 3 (bases 1 to 226979)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Worley, K.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished
                                                                                                                                                                                                                                                                                    Direct Submission
                                                                                                                                                                                                                                                                                                                 Rat Genome Sequencing Consortium.
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hgsc.bcm.tmc.edu/projects/rat/). As a result, the extend beyond the ends of the clone and there may be
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                                                                                                                                                                                                                                                                                                                                                                                    TX 77030,
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lor College
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;

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FEATURES
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                     AC096132
AC096132.6 GI:30522660
HTG; HTGS_PHASE1; HTGS_DRAFT; HTGS_ENRICHED.
RATTUS norvegicus (Norway rat)
Rattus norvegicus
Eukaryota;
                                                                                                                                     ***, 2 unordered pieces.
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                                                                                                                                                                                     AC096132
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                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
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NOTE: This is a 'working draft' sequence. It currently consists of 2 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 54404 a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence as soon as it is available and the accession number will be preserved.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Assembly program: Phrap; version 0.990329
Consensus quality: 196911 bases at least Q40
Consensus quality: 199136 bases at least Q30
Consensus quality: 200537 bases at least Q30
Consensus quality: 200537 bases at least Q20
Estimated insert size: 209329; sum-of-contigs estimation
Quality coverage: 6x in Q20 bases; sum-of-contigs estimation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: hgsc-help@bcm.tmc.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Center project name: GYAP Center clone name: CH230-63023
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                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note="clone_boundary
clone_end:T7
site:EcoRI
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clone_end:Sp6
site:EcoRI
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/note="wgs_contig"
complement(223202.
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/mol_type="genomic DNA"
/db_xref="taxon:10116"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nd_sequence:BH321164"
50080 c 47417 g 49761 t 25317 others
Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sequence:BH321166"
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                                                                                                                                                                                   231014 bp
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Allen, I.V., Aboysel, N., Ayodeji, M., Baca, E., Badden, H.,
Baldwin, D., Bandaranalke, D., Barber, M., Barrascad, M., Benahmed, F.,
Bladwin, D., Bandaranalke, D., Barber, M., Barrascad, M., Benahmed, F.,
Bladwald, K., Blair, J., Blankenburg, K., Blyth, P., Berom, M.,
Bryant, M., Buhay, C., Burch, P., Burrell, K., Calderon, E.,
Chacko, J., Chavez, D., Chen, G., Chen, R., Chen, Y., Chen, Z., Chu, J.,
Chacko, J., Chavez, D., Chen, G., Chen, R., Chen, Y., Chen, Z., Chu, J.,
Chaveland, C., Cookrell, R., Cox, C., Coyle, M., Cree, A., D'Souza, L.,
Davila, M.L., Davis, C., Day-Carroll, L., De Anda, C., Dederich, D.,
Delgado, O., Denson, S., Deramo, C., Ding, Y., Dinh, H., Divya, K.,
Eggn, A., Escotto, M., Bugene, C., Evans, C.A., Falls, T., Fan, G.,
Fraser, C., M., Gabisi, A., Ganta, R., Garrer, M., Guevara, W.,
Gunaratne, P., Haaland, W., Hamil, C., Hamilton, K.,
Harvey, Y., Havlak, P., Hawes, A., Henderson, N., Hernandez, M.,
Hernandez, R., Hines, S., Hladun, S.L., Hodgson, N., Hornandez, M.,
Hernandez, R., Hines, S., Hladun, S.L., Hodgson, A., Hogues, M.,
Hennins, B., Howells, S., Hulyk, S., kume, J., Idebind, D., Jackson, L., Jacob, L., Jiag, H., Johnson, B., Johnson, R., Johnson, R.,
Karpathy, S., Kaft, C.L., Lebow, H., Levan, J., Lowis, L., Li, Z., Liu, J.,
Liu, J., Liu, W., Liu, Y., London, P., Longacre, S., Lopez, J.,
Karpathy, S., Kelly, S., Khan, Z., King, L., Kora, C.,
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Mahleshwari, M., Mahindartne, M., Mahmoud, M., Malloy, K., Manjum, A.,
Maheshwari, M., Mahindartne, M., Mahmoud, M., Malloy, K., Manjum, A.,
Maheshwari, M., Mahindartne, M., Mahmoud, M., Malloy, K., Marjum, A.,
Mankervis, C., Nainer, G., Minja, E., Montemayor, Ju., Moore, S.,
Milosavjeric, A., Miner, G., Minja, E., Montemayor, Ju., Moore, S.,
Milosavjeric, A., Mortis, S., Minja, E., Montemayor, J., Moore, S.,
Milosavjeric, A., Soherer, S., Scott, G., Shateman, S., Shen, H.,
Shetty, J., Shartenbeyn, A., Sisson, I., Siteer
                                                                                                                                                                         Submitted (10-MAY-2003) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA On May 10, 2003 this sequence version replaced gi:24818180. The sequence in this assembly is a combination of BAC based reads
                                                                                                                                                                                                                                                                                                                                                                                                                            of Molecular and Human Genetics, Baylor
Baylor Plaza, Houston, TX 77030, USA
3 (bases 1 to 231014)
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                             and whole genome shotgun sequencing reads assembled using Atla (http://www.hgsc.bcm.tmc.edu/projects/rat/). Each contig descrin the feature table below represents a scaffold in the Atlas assembly (a 'contig-scaffold'). Within each contig-scaffold,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Submitted (17-SEP-2001) Human Genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Worley, K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unpublished
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contigs are ordered and oriented,
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                                                                                                          described
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FEATURES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             by sized gaps filled with Ns to the estimated size. The sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             shotgun sequence only contigs will be indicated in the feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NOTE: Estimated insert size may differ from sequence length (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.)
NOTE: This is a 'working draft' sequence. It currently consists of 2 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence.
                                                                                                Similarity
GGCTGGTGTCACCTGTT 122064
                                     GGCTGGTGTCACCTGTT 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         as soon as it is available and the accession number will be preserved.
                                                                                                                                                                               61558
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Center clone name: CH230-11C18

Center clone name: CH230-11C18

Center clone name: CH230-11C18

Center Summary Statistics

Assembly program: Atlas 3.0;

Consensus quality: 208633 bases at least Q40

Consensus quality: 212147 bases at least Q30

Consensus quality: 2121886 bases at least Q20

Estimated insert size: 214070; sum-of-contigs estimation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Quality coverage: 6x in Q20 bases; sum-of-contigs estimation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: hgsc-help@bcm.tmc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Web site:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Center code: BCM
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                                                                                                                                                                                                                                                                              /note="wgs_contig"
complement(228041.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               clone_end:T
                                                                                                                                                                                                                                                                                                                      note="wgs_contig"
169881. .171027
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        note="clone_boundary:
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clone end:T7"
                                                                                                                                                                                                                                                             'note="clone
                                                                                                                                                                                                                                                                                                                                                             note="wgs_contig"
24270. .126190
                                                                                                                                                                                                                                                                                                                                                                                                     'note="wgs_contig"
.21675. .123616
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               clone="CH230-11C18"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        organism="Rattus norvegicus"
                                                                                                                                                                           nd_sequence:BH320830"
48903 c 47136 g 58505 t
                                                                                                                                                                                                                                        .one_end:Sp6
                                                                     94.4%; Pr
100.0%; Pr
0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          sequence: BH320811"
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_xref="taxon:10116"
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                                                                                            Score 17; DB 2; pred. No. 2.9;
                                                                                                                                                                                                                                                           boundary
                                                                             Mismatches
                                                                                                                                                                                                                                                                                  .228684)
                                                                                                                                                                               14912 others
                                                                                                                     Length 231014;
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                                                                                                FEATURES
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                                                                               source
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2330
28155
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Chemistry: Dye-terminator Big Dye; 100% of reads Assembly program: Phrap; version 0.990319
Consensus quality: 236489 bases at least Q40
Consensus quality: 236478 bases at least Q30
Consensus quality: 237189 bases at least Q20
Insert size: 183000; agarcse-fp
Insert size: 248898; sum-of-contigs
Quality coverage: 16.05 in Q20 bases; agarcse-fp
Quality coverage: 12.12 in Q20 bases; sum-of-cont
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Submitted (23-FEB-2003) Genome Sequencing Center, 4444 Forest Parkway, St. Louis, MO 63108, USA 3 (bases 1 to 238552) McPherson,J.D. and Waterston,R.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Center: Washington University Genome Sequencing Center Center code: WUGSC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Submitted (01-MAR-2003) Genome Sequencing Center, 4444 Forest Parkway, St. Louis, MO 63108, USA
On Mar 1, 2003 this sequence version replaced gi:28475433.
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McPherson, J.D. and Wat
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McPherson, J.D. and Waterston, R.H.
The sequence of Mus musculus clone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequencing vector: M13; 0%
Sequencing vector: plasmid; 100%
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HTG; HTGS_PHASE1;
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Mus musculus chromosome UNK clone RP23-255C4, WORKING DRAFT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chemistry: Dye-primer ET; 0% of reads
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: submissions@watson.wustl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Web site:http://genome.wustl.edu/gsc/index.shtml
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE, 4 unordered pieces.
                                                                                                                                                                                                                                                                                                                                                                                     arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence as soon as it is available and the accession number will be preserved.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NOTE: This is a 'working draft' sequence. It currently consists of 4 contigs. The true order of the pieces is not known and their order in this sequence record is
/mol_type="genomic DNA"
/db_xref="taxon:10090"
                                                             organism="Mus musculus"
                                                                                                                                Location/Qualifiers
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                                                                                                     .238552
                                                                                                                                                           2229: contig of 2229 bp in length
2322: gap of unknown length
28154: contig of 25825 bp in length
28254: gap of unknown length
98428: contig of 70174 bp in length
98428: gap of unknown length
98528: gap of unknown length
238552: contig of 140024 bp in length
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  Allen,C., Allen,H., Alsbrooks,S., Amin,A., Anguiano,D.,
Allen,C., Allen,H., Alsbrooks,S., Amin,A., Anguiano,D.,
Anyalebechi, W., Aoyagi,A., Ayoegi,M., Baca,E., Baden,H.,
Baldwin,D., Bapdaranaike,D., Barber,M., Barnstead,M., Benahmed,F.,
Biswalo,K., Blair,J., Blankenburg,K., Blyth,P., Brown,M.,
Bryant,N., Buhay,C., Burch,P., Burrell,K., Calderon,E.,
Chacko,J., Chavez,D., Chen,G., Chen,R., Chen,Y., Chen,Z., Chu,J.,
Chacko,J., Chavez,D., Chen,G., Chen,R., Chen,Y., Chen,Z., Chu,J.,
Davila,M.L., Davis,C., Davy-Carroll,L., De Anda,C., Dederich,D.,
Delgado,O., Denson,S., Deramo,C., Ding,Y., Dinh,H., Divya,K.,
Draper,H., Dugan-Rocha,S., Dunn,A., Durbin,K., Duval,B., Eavee,K.,
Egan,A., Escotto,M., Eugene,C., Evans,C.A., Falls T., Fan,G.,
Fraser,C.M., Gabisi,A., Ganta,R., Garcia,A., Garner,T., Garza,M.,
Gebregeorgis,E., Geer,K., Gill,R., Grady,M., Guerra,W.,
Gunaratne,P., Haaland,W., Hamil,C., Hamilton,C., Hamilton,K.,
Hernandez,R., Hines,S., Hladun,S.L., Hodgson,A., Heyuevara,W.,
Gunaratne,P., Halland,W., Hamil,C., Hamilton,C., Hamilton,K.,
Harvey,Y., Havlak,P., Hawes,A., Henderson,N., Hernandez,J.,
Hernandez,R., Liu,Y., London,P., Longacre,S., Lopez,J.,
Liu,J., Liu,W., Liu,Y., Lowan,J., Lewis,L., Li,Z., Liu,J.,
Karpathy,S., Kelly,S., Kelly,S., Khan,Z., King,L., Koyar,C.,
Naheshwari,M., Mahindarne,M., Mahmoud,M., Malloy,K., Mangum,A.,
Mahshwari,M., Mahrer,G., Minja,E., Kontemayor,J., Moore,S.,
Mawhiney,S., Mel,D., Martin,K., Martin,R., Martin,R., Martin,R.,
Nankervis,C., Neal,D., Newton,N., Myuyen,N., Noris,S.,
Nankervis,C., Neal,D., Newton,N., Sisson,A., Pal,S., Parks,K.,
Palzon,M., Sison,I., Johnson,A., Reigh,R.,
Reilly,M., Reilly,M., Reny,Y., Reuter,M., Reiger,M.A., Reigh,R.,
Reilly,M., Sovery,G., Scherer,S., Scott,G., Shatsman,S., Shen,H.,
Shetty,J., Shvatrisbeyn,A., Sisson,I., Sitter,C.D., Smajs,D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AC131614
AC131614.3 GI:25138981
HTG; HTGS_PHASE1; HTGS_DRAFT; HTGS_FULLTOP.
Rattus norvegicus (Ngrway rat)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AC131614 244424 bp DNA linear Rattus norvegicus clone CH230-284C9, WORKING DRAFT unordered pieces.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            69322 a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (bases 1 to 244424)
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98529. 238552
/note="assembly_name:Contig26"
1 53309 c 49714 g 65901 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note="assembly_name:Contig22"
2330. 28154
/note="assembly_name:Contig24"
28255. .98428
Shvartsbeyn, A., Sisson, I.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     94.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
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FEATURES

235353 235453 240269

235352: contig of 235352 bp in length 235452: gap of unknown length 240268: contig of 4816 bp in length 240368: gap of unknown length 2404424: contig of 4056 bp in length.

240369

Location/Qualifiers

organism="Rattus norvegicus"

\* NOTE: Estimated insert size may differ from sequence length

(see http://www.hgsc.bcm.tmc.edu/docs/Genbank\_draft\_data.html).

\* NOTE: This is a "working draft' sequence. It currently

\* Consists of 3 contigs. The true order of the pieces

\* consists of 3 contigs. The true order of the pieces

\* a rout known and their order in this sequence record is

\* arbitrary. Gaps between the contigs are represented as

\* runs of N, but the exact sizes of the gaps are unknown.

\* This record will be updated with the finished sequence

\* This record will be updated with the finished sequence

Quality coverage: 7x in Q20 bases; sum-of-contigs estimation

as soon as it is available and the accession number will be preserved.

source

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AL Submitted (20-NOV-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
On Nov 20, 2002 this sequence version replaced gi:23195985.
On Nov 20, 2002 this sequence version replaced gi:23195985.
The sequence in this assembly is a combination of BAC based reads and whole genome shotgun sequencing reads assembled using Atlas (http://www.hgsc.bcm.tmc.edu/projects/rat/). Each contig described in the feature table below represents a scaffold in the Atlas assembly (a 'contig-scaffold'). Within each contig-scaffold, individual sequence contigs are ordered and oriented, and separated by sized gaps filled with Ns to the estimated size. The sequence contigs within a contig-scaffold that consist entirely of whole genome shotgun sequence reads. Both end sequences and whole genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sneed, A., Sodergren, E., Song, X.-Z., Sorelle, R., Sosa, J., Steinle, M., Strong, R., Sutton, A., Svatek, A., Tabor, P., Taylor, C., Taylor, T., Thomas, N., Thomas, N., Tingey, A., Trijos, Z., Usmani, K., Valas, R., Vera, V., Villasana, D., Waldron, L., Walker, B., Wang, J., Wang, Q., Wang, S., Warren, J., Warren, R., Weil, X., White, F., Williams, G., Willson, R., Weezyk, R., Wooden, H., Worley, K., Williams, G., Willson, R., Wleczyk, R., Wooden, H., Worley, K., Wright, D., Wright, R., Wu, J., Yakub, S., Yen, J., Yoon, L., Yoon, V., Wr, F., Zhang, J., Zhou, J., Zhou, X., Zhao, S., Dunn, D., Yon Niederhausern, A., Weise, R., Smith, D.R., Holt, R.A., Smith, H.O., Weinstock, G. and Gibbs, R.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Submitted (25-AUG-2002) Human Genome Sec
of Molecular and Human Genetics, Baylor
Baylor Plaza, Houston, TX 77030, USA
3 (bases 1 to 244424)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rat Genome Sequencing Consortium.
Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              shotgun
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Center clone name: CH230-284C9
------ Summary Statistics
Assembly program: Phrap; version 0.990329
Consensus quality: 221332 bases at least Q40
Consensus quality: 224113 bases at least Q30
Consensus quality: 224113 bases at least Q20
Estimated insert size: 226659; sum-of-contigs estimation
                                                                                                                                                                                                                                                                                                                                             Contact: hgsc-help@bcm.tmc.edu
                                                                                                                                                                                                                                                                                                  Center project name: KCCL
                                                                                                                                                                                                                                                                                                                                                                                                                     Web site: http://www.hgsc.bcm.tmc.edu/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Center: Baylor College
Center code: BCM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sequence only
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Genome
                                                                                                                                                                                                                                                                                                                                        Project Information
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              contigs will be indicated in the feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          of Medicine
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REFERENCE AUTHORS TITLE

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                KEYWORDS
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Bryant, N., Bulay, C., Buffel, F., Buffell, K., Carderon, E., Cardenas, V., Carter, K., Cavazos, I., Ceasar, H., Center, A., Chacko, J., Chavez, D., Chen, G., Chen, R., Chen, Y., Chen, Z., Chu, J., Cleveland, C., Cockrell, R., Cox. C., Coyle, M., Cree, A., D'Souza, L., Davila, M.L., Davis, C., Davy-Carroll, L., De Anda, C., Dederich, D., Delgado, O., Denson, S., Deramo, C., Ding, Y., Dinh, H., Divya, K., Delgado, O., Denson, S., Deramo, C., Ding, Y., Dinh, H., Divya, K., Delgar, H., Dugan-Rocha, S., Dunn, A., Durbin, K., Duval, B., Eaves, K., Delgar, H., Digar-Rocha, S., Dunn, A., Durbin, K., Duval, B., Eaves, K., Delgar, H., Dister, M., Foster, P., Fernandez, S., Finley, M., Flagg, N., Forbes, L., Foster, M., Foster, P., Fraser, C.M., Gabisi, A., Gall, R., Garcia, A., Garner, T., Garza, M., Gubregeorgis, E., Geet, K., Gill, R., Grady, M., Guerra, W., Guevara, W., Gunaratne, P., Haaland, W., Hamil, C., Hamilton, C., Hamilton, K., Harvey, Y., Havlak, P., Hawes, A., Henderson, N., Hernandez, J., Harvey, Y., Havlak, P., Hawes, A., Henderson, N., Hernandez, J., Hollins, B., Howells, S., Hladun, S.L., Hodgson, A., Hogues, M., Jackson, L., Jackson, E., Johnson, R., Li, Z., Liu, J., Liu, 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Muzny, D. Marie., Metzker, M. Lee., Abramzon, S., Adams, C., Alder, J., Allen, C., Allen, H., Alsbrooks, S., Amin, A., Anguiano, D., Anyalebechi, V., Aoyagi, A., Ayodegi, M., Baca, E., Baden, H., Baldwin, D., Bandaranaike, D., Barber, M., Barnstead, M., Benahmed, F. Biswalo, K., Blair, J., Blankenburg, K., Blyth, P., Brown, M., Bryant, N., Buhay, C., Burch, P., Burrell, K., Calderon, E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HTG; HTGS_PHASE2; HTGS_DRAFT; HTGS_ENRICHED RAttus norvegicus (Norway rat)
Rattus norvegicus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AC123215.3
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GCTGGTGTCACCTGTTA 203113
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clone_end:T7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note="wgs_end_extension
clone_end:T7"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="wgs_end_extension clone_end:T7"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          complement (3470.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nd_sequence:RXAHE17TJB"
54386 c 56712 g 59935 t 17274 others
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GI:23265875
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA linear HTG 21-SEP-2002
7, *** SEQUENCING IN PROGRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Barnstead, M., Benahmed, F.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
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                                                                                                                                                                                                                                                              FEATURES
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JOURNAL
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TITLE
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                              misc_feature
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Sanders, W., Savery, G., Scherer, S., Scott, G., Shatsman, S., Shen, H.,
Shetty, J., Shvartsbeyn, A., Sisson, I., Sitter, C.D., Smajs, D.,
Sneed, A., Sodergren, E., Song, X.-Z., Sorelle, R., Sosa, J.,
Steimle, W., Strong, R., Sutton, A., Svatek, A., Tabor, P., Taylor, C.,
Taylor, T., Thomas, N., Thomas, S., Tingey, A., Trejos, Z., Usmani, K.,
Valas, R., Vera, V., Villasana, D., Waldron, L., Walker, B., Wang, J.,
Wang, S., Warren, J., Warren, R., Wei, X., White, F.,
Williams, G., Willson, R., Wleczyk, R., Wooden, H., Worley, K.,
Wright, D., Wright, R., Wu, J., Yakub, S., Yen, J., Yoon, L., Yoon, V.,
Yu, F., Zhang, J., Zhou, J., Zhou, X., Zhao, S., Dunn, D., von
Niederhausern, A., Weiss, R., Smith, D.R., Holt, R.A., Smith, H.O.,
Weinstock, G. and Gibbs, R.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (21-SEP-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA On Sep 21, 2002 this sequence version replaced gi:21744493. The sequence in this assembly is a combination of BAC based reads and whole genome shotgun sequening reads assembled using Atlas (http://www.hgsc.bcm.tmc.edu/projects/rat/). As a result, the sequence may extend beyond the ends of the clone and there may be contigs that consist entirely of whole genome shotgun sequence reads. Both end sequences and whole genome shotgun sequence only contigs will be indicated in the feature table.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Submitted (29-MAY-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA 3 (bases 1 to 244636)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rat Genome Sequencing Consortium.
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Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                   NOTE: Estimated insert size may differ from sequence length (see http://www.hgsc.bcm.tmc.edu/docs/Genbank draft data.) NOTE: This is a 'working draft' sequence. It currently consists of 1 contigs. Gaps between the contigs are represented as runs of N. The order of the pieces is believed to be correct as given, however the sizes of the gaps between them are based on estimates that have provided by the submittor.
                                                                                                                                                                                                                                                      This sequence will be replaced
by the finished sequence as soon as it is available and
the accession number will be preserved.
1 244636: contig of 244636 bp in length.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Assembly program: Phrap; version 0.990329
Consensus quality: 223663 bases at least Q40
Consensus quality: 225326 bases at least Q40
Consensus quality: 225326 bases at least Q20
Consensus quality: 226436 bases at least Q20
Estimated insert size: 241958; sum-of-contigs estimation
Quality coverage: 4x in Q20 bases; sum-of-contigs estimation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Center project name: GSCP
Center clone name: CH230-118B7
Center Summary Statistics
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Web site: http://www.hgsc.bcm.tmc.edu/Contact: hgsc-help@bcm.tmc.edu
/note="clone_boundary
                                                                                                               /mol_type="genomic DNA"
/db_xref="taxon:10116"
                                                                                                                                                                                                                           1. .244636
                                                                     clone="CH230-118B7"
                                                                                                                                                                                     organism="Rattus norvegicus"
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COMMENT

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DEFINITION
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ORIGIN
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                                                                           Cardenas, V., Carter, K., Cavazos, I., Ceasar, H., Center, A., Cardenas, V., Carter, K., Cavazos, I., Ceasar, H., Center, A., Chen, Z., Chu, J., Clavez, D., Cchen, G., Cchen, R., Chen, Y., Chen, Z., Chu, J., Clavez, M., L., Davis, C., Davy-Carroll, L., De Anda, C., Dederich, D., Davis, C., Davy-Carroll, L., Debin, K., Duval, B., Eaves, K., Delgado, O., Denson, S., Denamo, C., Ding, Y., Dinh, H., Divya, K., Delgado, O., Denson, S., Dunn, A., Duvbin, K., Duval, B., Eaves, K., Egan, A., Escotto, M., Eugene, C., Evans, C.A., Falls, T., Fan, G., Egan, A., Escotto, M., Eugene, C., Evans, C.A., Falls, T., Fan, G., Farser, C. M., Gabisi, A., Ganta, R., Gardy, M., Gerrar, M., Gebregoergis, E., Geer, K., Gill, R., Gardy, M., Guerra, W., Guvara, W., Gebregoergis, E., Gester, K., Gill, R., Gardy, M., Guerra, W., Guvara, W., Gebregoergis, E., Gester, K., Gill, R., Gardy, M., Guerra, W., Guvara, W., Gebregoergis, E., Gester, K., Gill, R., Grady, M., Guerra, W., Guvara, W., Gebregoergis, E., Gester, K., Gill, R., Grady, M., Guerra, W., Guvara, W., Gebregoergis, E., Gester, K., Gill, R., Grady, M., Guerra, W., Guvara, W., Gebregoergis, E., Gester, K., Hamilton, C., Hamilton, C., Hamilton, K., Harrondez, S., Huden, S., Hume, J., Idlebird, D., Jackson, A., Hollins, B., Howells, S., Hulyk, S., Hume, J., Idlebird, D., Jackson, A., Jolivet, A., Kallyk, S., Hume, J., Idlebird, D., Jackson, A., Levis, J., Li, Z., Liu, J., Lorensuhewa, L., Li, Z., Liu, J., Lorensuhewa, L., Li, Z., Liu, J., Lorensuhewa, L., London, P., Lorgacre, S., Lopez, J., Liu, J., Lorensuhewa, L., London, P., Lorgacre, S., Lopez, J., Liu, J., Lorensuhewa, L., Lu, Y., Martin, K., Martin, R., Martinez, E., Moltemayor, J., Moore, S., Mangum, A., Mahmodu, M., Malloy, K., Mangum, A., Martin, K., Martin, R., Martinez, E., Mortensyor, J., Moore, S., Martin, R., Pal, S., Parks, K., Paster, R., Ruiz, S., Parks, K., Paster, R., Ruiz,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HTG; HTGS_PHASE1; HTGS_DRAFT; HTGS_FULLTOP.
Rattus norvegicus (Norway rat)
Rattus norvegicus
Rattus norvegicus
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Bukaryota; Metazoa; Chordata; Sciurognathi; Muridae; Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 Similarity
17; Conserv
Sanders,W., Savery,G., Scherer,S., Scott,G., Shatsman,S., Shen,H.
Shetty,J., Shvartsbeyn,A., Sisson,I., Sitter,C.D., Smajs,D.,
Sneed,A., Sodergren,E., Song,X.-Z., Sorelle,R., Sosa,J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Baldwin,D., Bandaranaike,D., Barber,M., Barnstead,M., Ben. Biswalo,K., Blair,J., Blankenburg,K., Blyth,P., Brown,M., Bryant,N., Buhay,C., Burch,P., Burrell,K., Calderon,E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Muzny,D.Marie., Metzker,M.Lee., Abramzon,S., Adams,C.,
Allen,C., Allen,H., Alsbrooks,S., Amin,A., Anguiano,D.
Anyalebechi,V., Aoyagi,A., Ayodeji,M., Baca,E., Baden,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              unordered pieces.
AC103080
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AC103080.6 GI:30579747
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complement(244009..2440)
/note="clone_boundary
clone_end:T7
site:EcoRI
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49128 c '49531 g 65
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Taylor, T., Thomas, N., Thomas, S., Tingey, A., Trejos, Z., Usmani, K., Valas, R., Vera, V., Villasana, D., Waldron, L., Walker, B., Wang, J., Wang, S., Warren, J., Warren, R., Wei, X., White, F., Williams, G., Willson, R., Wleczyk, R., Wooden, H., Worley, K., Wright, D., Wright, R., Wu, J., Yakub, S., Yen, J., Yoon, L., Yoon, V., Yu, F., Zhang, J., Zhou, X., Zhao, S., Dunn, D., von Niederhausern, A., Weiss, R., Smith, D.R., Holt, R.A., Smith, H.O., Direct Submission
                               and whole genome shotgun sequencing reads assembled using Atlas (http://www.hgsc.bcm.tmc.edu/projects/rat/). Each contig described in the feature table below represents a scaffold in the Atlas assembly (a 'contig-scaffold'). Within each contig-scaffold, individual sequence contigs are ordered and oriented, and separated by sized gaps filled with Ns to the estimated size. The sequence may extend beyond the ends of the clone and there may be sequence contigs within a contig-scaffold that consist entirely of whole genome shotgun sequence reads. Both end sequences and whole genome shotgun sequence reads. Both end sequences and whole genome shotgun sequence reads.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
On May 13, 2003 this sequence version replaced gi:23123516. The sequence in this assembly is a combination of BAC based reads
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Submitted (24-NOV-2001) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Submitted (13-MAY-2003) Human Genome Sequencing Center, Depa of Molecular and Human Genetics, Baylor College of Medicine,
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Direct Submission
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Genome Center
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AUTHORS

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Center project name: GJEC

Center clone name: GJEC

Center clone name: GH20-22501

Center clone name: CH230-22501

Center clone name: CH230-22501

Center clone name: CH230-22501

Consensus quality: 213558 bases at least Q40

Centersus quality: 215685 bases at least Q40

Centersus quality: 215685 bases at least Q20

Centersus quality: 217422 bases at least Q20

Estimated insert size: 218947; sum-of-centigs estimation

Quality coverage: 8x in Q20 bases; sum-of-centigs estimation Web site: http://www.hgsc.bcm.tmc.edu/Contact: hgsc-help@bcm.tmc.edu Center: Baylor C Project Information College of Medicine

NOTE: Estimated insert size may differ from sequence length
(see http://www.hgsc.bcm.tmc.edu/docs/Genbank draft data.html).

NOTE: This is a 'working draft' sequence. It currently
consists of 5 contigs. The true order of the pieces
is not known and their order in this sequence record is
arbitrary. Gaps between the contigs are represented as
runs of N, but the exact sizes of the gaps are unknown.
This record will be updated with the finished sequence
as soon as it is available and the accession number will be preserved 243313 243413 244624 244724 245934 243312: contig of 243312 bp in length
243412: gap of unknown length
244623: contig of 1211 bp in length
244723: gap of unknown length
245933: contig of 1210 bp in length
24633: gap of unknown length
247253: contig of 1220 bp in length
247253: gap of unknown length
247253: gap of unknown length .248793 248793: contig of 0 bp in length
n length
0 bp in length.

source

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1 (bases 1 to 50)
Edwards, C.A., Cantor, C.R., Andrews, B.M., Turin
Method of determining DNA sequence preference
molecule.
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Sorge, J.A. and Mullinax, R.L.
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Edwards,C.A., Cantor,C.R., Andrews,B.M., Turin,L.M. and Fry,K.E.
Sequence directed DNA binding molecules compositions and methods
Patent: US 6384208-A 546 07-MAY-2002;
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(bases 1 to 50)

Edwards, C. R., Cantor, C. R., Andrews, B.

Edwards, C. R., Cantor, C. R., Andrews, B.

Sequence-directed DNA-binding molecul

Patent: US 5578444-A 546 26-NOV-1996;
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1 (bases 1 to 435)

Edwards, J. B.D.M., Duclair, E. and Jordan, J.Y.

Sequence tag and encoded human protein:

Patent: JP 2001269182-A 1767 02-OCT-2001;
                                              16;
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Edwards,C.A., Cantor,C.R., Andrews,B.M. and Turin,L.M.
Edwards,C.A., Cantor,C.R., Andrews,B.M. and Turin,L.M.
Screening assay for the detection of DNA-binding molecules
Patent: US 5726014-A 546 10-MAR-1998;
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GGCTGGTGTCACCTGT 171
                       GGCTGGTGTCACCTGT 16
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JP 2001269182-A/1767
02-OCT-2001
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26-FEB-1999 US 60/122487
JEAN BAPUTIST DUMAS MILNE EDWARDS, EIMERIC DUCLAIR, JEAN YVES
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14 c 10 g
                                                                                                    /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xmef="taxon:9606"
114 c 133 g 7
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Location/Qualifiers
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Direct Submission

Submitted (13-MAY-2003) BD Biosciences Clontech, 1020 East Meadow Circle, Palo Alto, CA 94303, USA

This CDS clone is a part of a collection of human full length expression clones generated by BD Biosciences Clontech and the Harvard Institute of Proteomics. Each CDS has been cloned in two forms: with and without stop-codon (to allow fusion with C-terminal tag). The CDS has been directionally cloned using BD In-Fusion(TM) cloning system between the SalI and HindIII sites of the pDNR-DUAL vector. Additional sequences in the clone: 'ACC' after SalI site and before 'ATG' to provide Kozak consensus sequence; 'GG' after last codon and before HindIII site to maintain reading frame. Clone distribution: http://bboinfo.clontech.com/orfclones.
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Sequence
AX321046
                                                                                                                                                                                                                                                                                                                                                                                                     1 (bases 1 to 963)
Kounding,N., Chen,X., Rolfs,A., Halleck,A., Hines,L., Eisenstein,S., Koundinga,M., Raphael,J., Moreira,D., Kelley,T., LaBaer,J., Lin,Y., Phelan,M. and Farmer,A.
Cloning of human full-length CDSs in BD Creator(TM) System Donor
                                                                                                                                                                                                                                                                                 2 (bases 1 to 963)
Kalnine, N., Chen, X., Rolfs, A., Halleck, A., Hines, L., Eisenstein, S., Koundinya, M., Raphael, J., Moreira, D., Kelley, T., LaBaer, J., Lin, Y., Phelan, M. and Farmer, A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BT006889.1 FLI_CDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens eukaryotic translation delta, 44kDa mRNA, complete cds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent: WO 0177168-A 63 18-OCT-2001; CORIXA CORPORATION (US)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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This CDS clone is a part of a collection of human full length expression clones generated by BD Biosciences Clontech and the Harvard Institute of Proteomics. Each CDS has been cloned in two forms: with and without stop-codon (to allow fusion with C-terminal tag). The CDS has been directionally cloned using BD In-Fusion(TM) cloning system between the Sall and HindIII sites of the pDNR-DUAL vector. Additional sequences in the clone: 'ACC' after Sall site and before 'ATG' to provide Kozak consensus sequence; 'GG' after last codon and before HindIII site to maintain reading frame. Clone distribution: http://bioinfo.clontech.com/orfclones.
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Kalnine, N., Chen, X., Rolfs, A., Halleck, A., Hines, L., Eisenstein, S., Koundinya, M., Raphael, J., Moreira, D., Kelley, T., LaBaer, J., Lin, Y., Phelan, M. and Farmer, A.
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1 (bases 1 to 963)

Kalnine, N., Chen, X., Rolfs, A., Halleck, A., Hines, L., Kalnine, M., Raphael, J., Moreira, D., Kelley, T., Labe Koundinya, M., Raphael, J., Moreira, D., Kelley, T., Labe Phelan, M., and Farmer, A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished
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/product="eukaryotic translation initiation subunit 4 delta, 44kDa"
/protein_id="AAP35535.1"
/db_xref="GI:30582617"
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291 c 288 g 147 t
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PELLPGAPLPPKEVINGNIKTVTEYKIDEDGKKFKIVRTFRIETRKASKAVARRKNW
KKFGNSEFDPPGPNVATTTVSDDVSMTFITSKEDLNCQEEEDPMNKLKGQKIVSCRIC
/mol_type="mRNA"
/db xref="taxon:32630"
/clone="db0457L1.0"
/clone_lib="BD Creator(TM) CDS Library derived from MGC
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/clone_lib="BD Creator(TM)
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LaBaer,J., Lin,Y.,
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Sequence 3 from Patent US 4707358.
104174
                                                                                                                 Homo sapiens eukaryotic translation initiation (p42) mRNA, complete cds.
AF020833
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Kieff, E., Tanner, J., Hummel, M. and Beisel, C.
Vaccine against Epstein-Barr Virus
Patent: US 4707358-A 3 17-NOV-1987;
The University of Chicago; Chicago, IL
Location/Qualifiers
          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 1103)
                                                       Homo sapiens (human)
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Bandyopadhyay, A. and Maitra, U.
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PELLPGAPLPPPKEVLNGNIKTVTEYKIDEDGKKFKIVRTFRIETRKASKAVARRKNW
KKFGNSEFDPPGPNVATTTVSDDVSMTFITSKEDLNCQEEEDPMNKLKGQKIVSCRIC
KGDHWTTRCPYKDTLOPMQKELAEQLGLSTGEKEKLPGELEPVQATQNKTGKYVPPSL
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291 c 288 g 148 t
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383 c 199 g
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factor 3, subunit 4 delta, 44kDa"
/protein id:"AAP36240.1"
/db_xref="GI:30583983"
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/transl_table=11
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/note="Vector: pDNR-Dual"
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U96074.1
                                                                           2 (bases 1 to 1115)
Block, K.L., Vornlocher, H.-P.
Direct Submission
                                                                                                                                                     1. (bases 1 to 1115)
Block, K.L., Vornlocher, H.P. and Hershey, J.W.
Characterization of cDWAs encoding the p44 and p35
human translation initiation factor eIF3
J. Biol. Chem. 273 (48), 31901-31908 (1998)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (26-AUG-1997) Dev. & College of Medicine of Yeshiva Avenue, Bronx, NY 10461, USA Location/Qualifiers
                                             Submitted (01-APR-1997) Biological Chemistry, U C Davis, Tupper Hall, Davis, CA 95616, USA
                                                                                                                                                                                                                                                                              Homo
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Human translation initiation
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                                                                                                                                                99041954
                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                          Homo sapiens (human)
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KGDHWTTRCPYKDTLGPMQKELAEQLGLSTGEKEKLPGELEPVQATQNKTGKYVPPSL
RDGASRRGESMQPNRRADDNATIRVTNLSEDTRETDLQELFRPFGSISRIYLAKDKTT
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PELLPGAPLPPPKEVINGNIKTVTEYKIDEDGKKFKIVRTFRIETRKASKAVARRKNW
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/db_xref="GI:2460200"
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mol_type="mRNA"
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University, 1300, Morris
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                                                                                                                                                                                               subunits
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ORGANISM
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BC000733/c
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VERSION
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TITLE
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Best Local
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                                                                                                                                                                                                                                                                                                                                                        cDNA Library Preparation: Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Institute for Systems Biology
                                                                                                                                                                                                                                                                                                                                                                                                       Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BC000733 1128 bp mRNA linear PRI 13 Homo sapiens, eukaryotic translation initiation factor 3, (delta, 44kD), clone MGC:2053 IMAGE:3504640, mRNA, complet
                                                                                                                                                                                                                                                                                                           http://www.systemsbiology.org
contact: amadan@systemsbiology.org
Anup Madan, Rachel Dickhoff, Jessi
                                                                                                                                                                                                                                                                                                                                                                                                                                       NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
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BC000733.1 GI:12653882
                                                                                                                                                                                                                                                                                            Greene, Mark Ketteman and Anuradha Madan
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ilarity 100.0%;
Conservative
                                                                                                              /db_xref="LocusID:8666"
                                                                                                                               Tom/
                                                                                                                                              organism="Homo
                                                                                                                                                                             Location/Qualifiers
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                                                                                              xref="taxon:9606"
                                                                                                                  type="mRNA"
ref=""
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Pred. No.
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Length 1115;
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complete cds. PRI 12-JUL-2001

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Gaps

0;

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 1128)

Submitted (15-NOV-2000) National Institutes of Health, Mammalian Gene Collection (MGC), Cancer Genomics Office, National Cancer Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,

Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Series: IRAL Plate: 7 Row: e Column: 11
This clone was selected for full length sequencing because it passed the following selection criteria: matched mRNA gi: 10280561 /clone="MGC:2053 IMAGE:3504640"
/tissue type="Placenta, choriocarcinoma"
/clone lib="NIH MGC 21"
/lab\_host="DH10B-R"
/lab\_host="DH10B-R"

BASE COUNT ORIGIN

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Query Match
Best Local Similarity
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This clone was selected for full length sequencing because it passed the following selection criteria: Similarity but not identify the contents of the collowing selection criteria:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                cDNA Library Preparation: CLONTECH Laboratories, Inc.
cDNA Library Agrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Sequencing Group at the Stanford Human (
Center, Stanford University School of Medicine, Stanford,
Web site: http://www-shgc.stanford.edu
Contact: (Dickson, Mark) mcd@paxil.stanford.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (25-MAY-2001) National Institutes of Health, Mammalian Gene Collection (MGC), Cancer Genomics Office, National Cancer Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 1138)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Dickson, M., Schmutz, J., Grimwood, J., Rodriquez, A., and Myers,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Direct Submission
                                                                                                                                                                                                                                                  identity to protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1138 bp mRNA linear PRI 12-JUL-2001 sapiens, eukaryotic translation initiation factor 3, subunit 4 sapiens, clone MGC:14741 IMAGE:4279770, mRNA, complete cds.
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1336 c 335 g 174 t
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KGDHWTTRCPYKDTLGPMQKELAEQLGLSTGEKEKLPGELEPVQATQNKTGKYVPPSL
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/tissue type="Brain, primitive ne
/clone_Tib="NIH_MGC_56"
/lab_host="DH10B"
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subunit 4 (delta, 44kD)"
                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                             organism="Homo sapiens"
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  note="Vector:
                                                                                                             xref="LocusID:8666"
xref="taxon:9606"
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Pred. No. 15;
0; Mismatches
pDNR-LIB
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AUTHORS
TITLE
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                                                                    RESULT 26
AF094850/c
LOCUS
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BD063236/c
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DEFINITION
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REMARK COMMENT

FEATURES

CDS

REFERENCE AUTHORS TITLE

JOURNAL

SOURCE ORGANISM

ACCESSION VERSION

KEYWORDS

RESULT 24 BC008469/c

DEFINITION Focus 밁

Matches

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LASSIFIED

L (bases 1 to 1142)

Escobedo, J., Hu,O., Garcia,.

Secreted human proteins

JOURNAL Patent: JP 2001505783-A 11 08-
CHIRON CORP

PM JP 2001505783-A/1*

PD 08-MAY-2001

PF 11-DEC-1°

PR 11-DEC-1°

PI 11-DEC-1°
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                           PI JAIME ESCOBEDO,QUIANJIN HU,PABLO GARCIA,LEWIS T WILLIAMS PI

SRINIVAS KOTHAKOTA

PC C12N15/12,C12N15/62,C12N15/85,C12N5/10,C12N1/21,C07K14/47, PC

C07K16/18

CC Topology: Linear;

FH Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BD063236
Secreted
                                              AF094850 1174 bp mRNA linear PRI Homo sapiens eukaryotic translation initiation factor 3
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JP 20015057
p42/p44 mRNA, complete
AF094850
AF094850.1 GI:10280561
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                                                                                                                                                                                                                              Similarity
                                                                                                                                              GGCTGGTGTCACCTGT 144
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08-MAY-2001
11-DEC-1997 JP 1998526977
11-DEC-1996 US 60/032757
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/protein_id="AAH08469.1"

/db_xref="G::14250114"

/translation="MPTGDFDSKPSWADQVEEEGEDDKCVTSELLKGIPLATGDTSPE
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KGDHWTTRCPYKDTLGPMQKELAEQLGLSTGEKEKLPGELEPVQATQNKTGKKVPPSL
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1 334 c 331 g 176 t
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                                                                                                                                                                                                                                                                                            /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
_335 c 331 g 17
                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /product="eukaryotic translation initiation factor 3,
subunit 4 (delta, 44kD)"
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                                                                                                                                                                                                             Score 16; DB; Pred. No. 15; 0; Mismatches
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or 3 subunit
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TITLE
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Best Local Similarity
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source
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  168 GGCTGGTGTCACCTGT 153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16;
                                                                                                                                                           Desmodesmus pirkollei
Desmodesmus pirkollei
Bukaryota; Viridiplantae; Chlorophyta; Chlorophyceae;
Chlorococcales; Scenedesmaceae; Desmodesmus.
1 (bases 1 to 2199)
Hegewald, E., Coesel, P. and Hegewald, P.
A phytoplankton collection from Bali, with description of a
               Hegewald, E., Coesel, P. and Hegewald, P. Direct Submission
Submitted (13-FEB-2001) Institute of Chemistry and Dyna Geosphere 6, Research Center Juelich, Juelich D-52425,
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 GGCTGGTGTCACCTGT 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Mammalia; Eutheria; Primates; Catarrhini 1 (bases 1 to 1174)
Chen, W., Blough, R.I. and Winkelmann, J.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Submitted (24-SEP-1998) Internal Medicine/Hematology-Oncology, University of Cincinnati College of Medicine, 231 Bethesda Ave., Cincinnati, OH 45267-0508, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Molecular cloning, genomic structure and chromosomal localization of a novel human RNA binding protein gene homologous to a tumor necrosis factor alpha inducible transcript in mouse
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AF348496.1
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                                                                                                                             Unpublished
                                                                                                                                              Desmodesmus species (Chlorophyta,
                                                                                                                                                                                                                                                                                                                                                  Desmodesmus pirkollei 18S
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                                                                                                           (bases 1 to 2199)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /db_xref="gi:10280562"
/translation="MPTGDFDSKPSWADQVEEEGEDDKCVTSELLKGIPLATGDTSPE
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KCHWATTRCPYKDTLGPMQKELAEQLGLSTGEKEKLPGELEPVQATQNKTGKYVPPSL
RCHWATTRCPYKDTLGPMQKELAEQLGLSTGEKEKLPGELEPVQATQNKTGKYVPPSL
RDGASRFISEFHREEDAARAIAGVSGFGYDHLILNVEMAKPSTN"

336 c 331 g 173 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /product="eukaryotic translation initiation
subunit p42/p44"
/protein_id="AAGI5419.1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        note="eIF3 p42/p44"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/chromosome="19"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 16;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                        2199
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                                                                                                                                                  Scenedesmaceae
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1 RNA gene, complete sequence.
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                                                     Dynamics
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AF172333/c
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 (bases 1 to 2556)
Chang,S.H., Kim,S.H., Lee,W.K., Kim,H.J., Choi,
Jang,H.S., Chung,G.H., Kwon,T.H., Kim,D.H., Yar
Cloning and analysis of the Epstein-Barr virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 GGCTGGTGTCACCTGT 16
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Jang, H.S., Chung, G.H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AF172333 2556 bp mRN
Human herpesvirus 4 cell-line SNU-1103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (27-JUL-1999) Faculty of Biological Sciences,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              B-lymphoblastoid cell lines from cancer patients
In Vitro Cell. Dev. Biol. Anim. 34 (2), 97-100 (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human herpesvirus 4 (Epstein-Barr virus)
Human herpesvirus 4
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Lee, W.K., Kim, S.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gammaherpesvirinae; Lymphocryptovirus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           glycoprotein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cells 8 (5),
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ALTMRSKKLPINVTTGEEQQVSLESVDVYFQDVFGTMWCHHAEMQNPVYLLPETVPI
IKMDNCNSTMITAVVRAQGLDVTLPLSLPTSAQDSNFSVETQMLGMEIDIECIMEDGEI
SQVLPGDNKFNITCSGYESHVPSGGILSTSPPARJIPITGTGYAYSLFLTFRPVSRFLG
NNSILYVYSGGGFKASGGNVCIQSNIVFSDEIPASQDMFTNTTDITYVCDNATYSVP
MVTSEDANSPNVTVTAFMAMPNNTETDFKCKWTLTSGTPSGCENISGAFASNRTFDIT
VSGLGTAPMILITRFATNATTTTHKVIFSKAPESTTNTTGFAAENTTTGLES
STHVPTNLTAFASTGFTVSTADVTSFTBAGTTSGAFSVTFPSBSBDGTESKAPDMTS
STHVPTNLTAFASTGFTVSTADVTSFTBAGTTSGAFSTSSWSLRPSSISETLSPST
PTSAVTTFTPNATSPTPANTSPTLGKTSPTSAVTTFTNATSFTVGETSPQ
ANTTNHTLGGTSSTPVVTSPPKNATSAVTTGQHNITSSSTSSMSLRPSSISETLSPST
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465 c 593 g 569 t
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/cell_line="SNU-1103"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Korea
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     585-593 (1998)
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lee, W.K., Kim, H.J., Choi, S.H., Park, J.H., Kwon, T.H., Kim, D.H., Yang, M.S. and Jang, Y.S.
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Chang, S.H., Kim, S.H., Lee, W.K., Kim, H.J., Choi, S.H., Park, J.H., Jang, H.S., Chung, G.H., Kwon, T.H., Kim, D.H., Yang, M.S. and Jang, Y.S.
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Chang,S.H., Kim,S.H., Lee,W.K., Kim,H.J., Choi,S.H., Park,J.H.,
Jang,H.S., Chung,G.H., Kwon,T.H., Kim,D.H., Yang,M.S. and Jang,Y.S.
Cloning and analysis of the Epstein-Barr virus glycoprotein 350
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Submitted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 B-lymphoblastoid cell lines from cancer patients
In Vitro Cell. Dev. Biol. Anim. 34 (2), 97-100 (1998)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Submitted (27-JUL-1999) Faculty of Biological Sciences, C. National University, 664-14, Dukjin-dong 1-Ka, Dukjin-Ku,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Direct Submission
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Lee,W.K., Kim,S.M., Sim,Y.S., Cho,S.G., Park,S.H., Kim,C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gammaherpesvirinae; Lymphocryptovirus.
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TEPTIDYGGDSTTPRTRYNATTYLPPSISSKLRPRWTFTSPPVTTAQATVPVPPTSQP
RFSNLSMLVLQWASLAVLTLLLLLVMADCAFRRNLSTSHTYTTPPYDDAETYV"
a 798 c 532 g 513 t
MVTSEDANS PNVTVTAFWAWPNNTETDFKCKWTLTSĞTPSGCENISGAFASNRTFDIT
VSGLCTAPKTLLI TRTAINATTTTHKVL FSKAPESTTTSSTLAVTGFAAPNTTTGLPS
STHYPTNLTAFYSTGFTYSTADVTSFPPAGTTSGAS PVTPRESERDNGTESKAPDMTS
PTSAVTTPTPNATSPTPAVTTPTPNATSPTLGKTSPTSAVTTPTPNATSPTPAVTTPT
                                                                                    KWDNCNSTNITAVVRAQGLDVTLPLSLFTSAQDSNFSVKTEMLGNEIDIECIMEDGEI
SQVLFGDNKFNITCSGYESHVESGGILTSTSFVATFIFGTGYAYSLRLTFRFVSRFLG
NNS1LYVFYSGNGFKASGGDYCIQSNIVFSDEIFASQDMFTNTDITYVCDNATYSVF
                                                                                                                                                                                /translation="meaallvcqytiqsliqltrddpgffnveilefpfypacnvcta
dvnatinfdvggkkhklnldfglltphtkavyqprgafggsenatnlfllellgagel
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                                                                                                                                                                                                                 /codon_start=1
/prodmct="major outer envelope glycoprotein gp350"
/protein_id="AAD51697.1"
/db_xref="GI:5802483"
/db_xref="GI:5802483"
                                                                                                                                                           ALTMRSKKLPINITAGEEQQVSLESVDVYFQDVFGTMWCHHAEMQNPVYLIPETVPYI
                                                                                                                                                                                                                                                                                                                                       /organism="Human herpesvirus
/mol type="mRNA"
/db xref="taxon:10376"
/cell_line="SNU-20"
                                                                                                                                                                                                                                                                                                                                                                                                                                                          location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Korea
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glycoprotein 350/220.
Human herpesvirus 4 (Epstein-Barr virus)
Human herpesvirus 4
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X67776
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2 (bases 1 to 2661)
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DVNATINED VGGKKHKLINLDFGLLTBHTKAVYOPRAFGGSENATNLFILELLGAGEL
ALTMESKKLEIN ITTGEBONGLESUNYFODVEGHMACHNAPNNO PVLI BETUPY I
KWDNCNSTNITAVVRAGGLDVTLFLSLPTSAODSNESVKTEMLGNEID IECIMEDGEI
SQULPGDNKRNITAVVRAGGLDVTLFLSLPTSAODSNESVKTEMLGNEID IECIMEDGEI
SQULPGDNKRNITGSGYESHVPSGGILTSTSPVATTIFGTAYASLRLTPRRVSREIG
NNSILYVFYSGMGPRASGGDXCIOSNIVFSDEIPASODMPNITEDITYVGDNATYSVP
MYTSEDANSPNVTVTAAWAWPNNTETDEKCKWTLTSGTPSGCENISGAFASNRTFDIT
VSGLGTAFKTLAINATTTTHKVIFSKAPESTTTSPTLNATSPTAARNTTFDIT
VSGLGTAFKTLAINATTTTHKVIFSKAPESTTSPTLATTSFPRANTSPPR
STHVPTNITAPASTGPTVSTADVTSPTPAGTTSGASPVTFSPSBERDMGTESKAPDWTS
STHVPTNITAPASTGPTVSTADVTSPTPAGTTSGASTSTAVTTFTNATSPTRARTSPPAVTTPT
PNATIPTLARASTGPTVSTADVTSPTPAGTTSGASTSTSNTSHMPLLTSAHPTGGENIT
QVTRASTSTHAVSTSSPARP FGTTSQASGPGNSSTSTKPGENVNTKGTPFRANTSPOA
ATSAVTTGQHNITSSGSTSMSLRPSSISETLSPSTSDNSTSHMPLLTSAHPTGGENIT
QVTRASTSTHAVSTGGKANTTSPENATSPTGGARTSTSTEPTDYGGDSTSTRYNATTY
PSGGKTAVPTTSTGGKANTTGGARTTSGASGFGNSTSTKPGENVNTTSTPTRTRXNATTY
PSGGKTAVPTTSTGGKANTTGGARTTSGASGFGNSTSTKPGENVNTTTBTRTRXNATTY
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LVMADCAFRRNLSTSHTYTTPPYDDAETYV"
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PSGQKTAVPTVTSTGGKANSTTGGKHTTGHGARTSTEPTTDYGGDSTTPRTRYNATTY
                  LPPSTSSKLRPRWTFTSPPVTTAQATVPVPPTSQPRFSNLSMLVLQWASLAVLTLLLL
LVMADCAFRRNLSTSHTYTTPPYDDAETYV"
a 847 c 542 g 522 t
                                                                                                                                                                                                                                                                                                                                                                                                                               /product="glycoprotein 350/220"
/protein_id="CAA47986.1"
/db_xref="GI:59164"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /mol_type="genomic DNA"
/strain="P3HR1"
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                                                                                                                                                                                                                                                                                                                                                                                                            db_xref="SWISS-PROT:Q07284"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted (04-JUL-1996) S.D.V. Pepper, Paterson Institute for Cancer Research, Molecular Biology, Wilmslow Rd, Withington, Manchester M20 9BX, UK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2 (bases 1 to Pepper, S.D.V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mackett,M., Cox,C., Pepper,S.D., Lees,J.F., Naylor,B.A., Wedderburn,N. and Arrand,J.R. Immunisation of common marmosets with vaccinia virus expressing Epstein-Barr virus (EBV) gp340 and challenge with EBV J. Med. Virol. 50 (3), 263-271 (1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BLLF1 gene; gp340.
Human herpesvirus 4 (Epstein-Barr virus)
Human herpesvirus 5.
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GNSSTSTKPGEVNVTKGTPPKNATSPQAPSGQKTAVPTVTSTGGKANSTTGGKHTTGH
GARTSTEPTTDYGDDSTTPRPRYNATTYLPPSTSSKLRPRWTFTSPPVTTAQATVPVP
                                                                                                                                                                                                                                                                                                                                          /t-anslation="MEAALLYCQYTIQSLIHLTGEDPGFENUEI PEFPPYPTCNVCTA
DVNVTINPDVGGKKHQLJDLFGQLTPHTKAVYQPGAJFGSESHATNIFLLELLGAGEL
ALTMRSKKLPINVTTGEEQQVSLESVUDYYGDDVGTMWCHHAEMONPVYLIPETYPYI
KWDNCNSTNITAVVRAQGLDVTLPLSLPTSAQDSNFSVKTQMLGNIEIDIECIMEDGEI
SQVLFGDNKFNITGSGYESHVPSGGILTSTSPVATPIPGTTGTTYGDDATYTSPL
NNSLLYVFYSGNGPKASGGDYCIQSNIVFSDEIPASQDMPTNTTDITYGDDATYSPL
MYTSEDANSPNVTVTAFWAWPNNTETDFKCKWTLTSGTPSGCENISGAFASNRTFDIT
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STHVPTNLTAPASTGPTVSTADVTSPTPAGTTSGASPVTPSPS PRDNGTESKAPDMTS
PTSAVTTPTPNGTSPTPAMTTPTPNATSPTLGKTSPTSAVTTPTPNATSPTPAVTTPT
                                                                                                                                                                                        PTSQPRFSNLSMLVLQWASLAVLTLLLLLVMADCAFRRNLSTSHTYTTPPYDDAETYV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /product="gp340"
/protein_id="CAA67558.1"
/db_xref="GI:2769560"
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mol_type="genomic DNA"
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PC C12
PC C12
CC St1
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CC sty
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CC hyr
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FH Key
FT CDS
                                                                                                                                                                                            HS4GP34QA 1210 bp DNA linear VRL 15-E Epstein-Barr virus glycoprotein 340 (BLLF1) and BLLF2 genes, complete cds's, and BLLFR3 gene, first exon.
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Eriotsuto,K., Jiaroomu,T., Mearii,F. and Kurisutofuaa,B. VACCINE TO EPSTAIN VAR VIRUS
Patent: JP 1985232094-A 1 18-NOV-1985;
The Epstein-Barr virus candidate vaccine antigen gp340/220 is highly conserved between virus types A and B Virology 195 (2), 578-586 (1993)
                                                                                                                               Human herpesvirus 4
                                                                                                                                               antigen; glycoprotein 340; gp340; viral antigen
Human herpesvirus 4 (Epstein-Barr virus)
                                                                               Gammaherpesvirinae; Lymphocryptovirus.
1 (bases 1 to 3210)
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                                                  Arrand, J.R.
                                                               Lees, J.F., Arrand, J.E., Pepper, S.D., Stewart, J.P., Mackett, M. and
                                                                                                                Viruses; dsDNA viruses, no RNA stage; Herpesviridae;
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30-JAN-1984 US 84 575352, 23-JUL-1984 US
SUTO KIIFU, JIEROOMU TANNAA, MEARII FUNMERU,
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hypothetical: No;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2N15/00, A61K39/245, C07H21/04, C07K13/00, C12N1/00//C12P21/02,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              b
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /product='gp 220/200'.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Human herpesvirus 4"
/mol type="genomic DNA"
/db xref="taxon:10376"
876 c 557 g 526 t
                                                                                                                                                                                GI:291519
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1. .2721
                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                           88.9%; Score 16; DB 6; Length 2721; 100.0%; Pred. No. 14;
                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ביבו טף UNA linear
Epstein-Barr virus gp 220/200.
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                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                               Indels
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PI
                                                                                                                                                                                                                                                  VRL 15-FEB-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       633558 PI
KURISUTOFUAA
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RESULT 34
A11128/c
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  ACCESSION
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PUBMED
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                                                                                                                                                                                                                                                                       Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CDS
                                                                                                                                                                   601
All128 3400 bp DNA linear PAT 07-DEC-1993 Synthetic nucleotide sequence of the leftward reading frame of the Bam L-fragment encoding gp 250/350. All128
                                                                                                                                                                                                                                                 16;
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                                                                                                                                                                                                        GGCTGGTGTCACCTGT 16
                                                                                                                                                                                                                                                                                                                                                            659
                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           QVTPASTSTHHVSTSSPAPRPGTTSQASGPGNSSTSTKPGEVNVTKGTPPKNATSPQA
PSGQKTAVPTVTSTGGKANSTTGGKHTTGHGARTSTEPTTDYGGDSTTPRTRYNATTY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STHVETNLTAPASTGETVSTADVTSTPAGTTISGASFVTPSSERDNGTESKADMTS
ETSAVTTPIPANTSPTBAVTTPTENATSPTLKTSFTSAVTSPTRANTSPTPAVTTSP
PNATI PTLGKTSPTSAVTTPTENATSPTVGETISPOANTTNHTLGGTSSTPVUTSPEXD
ATSAVTTGQHNITSSSTSSMSLRPSSISETLSPSTSDNSTSHMPLLTSAHPTGGENIT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NNSILYVFYSGNGPKASGGDYCIQSNIVFSDEIPASQDMPTNTTDITYVGDNATYSVP
MVTSEDANSPNVTVTAFWAWPNNTETDFKCKWTLTSGTPSGCENISGAFASNRTFDIT
VSGLGTAPKTLIITRTATNATTTTHKVIFSKAPESTTTSPTLNTTGFAAPNTTTGLPS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /translation="Meaall/CoytigsLigitrdDpgffnveilefppypacnvcta
DVNATINFDVGGKCHKUNLDFGLLTPHTKAVYQPGAFGGSENATNIFILELIGAGEL
ALTMRSKKLPINITTGEEQQVSLESUDVYFQDVGTMWCHHAEMQNPVLILPETVYBL
KWDNCNSTNITAVVRAQGLDVTLPLSLPTSAQDSNFSVKTEMLGNEIDIECIMEDGEI
                                                                                                                                                                                                                                                                                                                                                                                                 2751. .3210
/gene="BLRF3"
                                                                                                                                                                                                                                                                                                                                                                                                                                                            /translation="MDKDRPGLPAPDDNIEEVPSTSGVQERASEGDWENVLIEISDSS
SEEEABDAHLDSSQRGKKRKKVDDDAGGSAPAQHVPPPQLDHPGREAILYRFPLDLRR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     complement (138. .584)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /codon_start=1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /trānslation="MCPPVRQRPAQAPPAKRQALETVPHPQNRGRLMSPKARPPKMQR
RPRPPVAKRRRFPRSPQQVERPILPPVESTPQDMEPGQVQSPPQITAVIQLRQERDTM
RPPIYLPALLANCGPAGLLRAHRLPQPKPPCLSRQRPSPDSQTSPC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              complement (138. .584)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SQVLPGDNKFNITCSGYESHVPSGGILTSTSPVATPIPGTGYAYSLRLTPRPVSRFLG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /product="glycoprotein 340"
/protein_id="AAA02783.1"
/db_xref="GI:291520"
                                                                                                                                                                                                                                                                                                                                                          /number=1
678 c
                                                                                                                                                                                                                                                                                                                                                                                                                                        FIQAIGAAATVSFPMAQVCDVCFCPSHNKVSDLLPLVSAPRHASHRPVFRI"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /protein_id="AAA02784.1"
/db_xref="GI:291521"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gene="BLLF2"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                'gene="BLLF1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         organism="Human herpesvirus 4"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .PPSTSSKLRPRWTFTSPPVTTAQATVPVPPTSQPRFSNLSMLVLQWASLAVLTLLLL
                                                                                                                                                                                                                                                                     88.9%;
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                                                                                                                                                                                                                                                                     Score 16; DB 14; Length 3210; Pred. No. 14;
                                                                                                                                                                                                                                                   Mismatches
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                                                          linear PAT 07-DEC-1993
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AUTHORS
TITLE
                                                                                                                                                                                                                            SOURCE
ORGANISM
                                                                                                                                                                                                                                                                                                            DEFINITION
ACCESSION
                                                                                                                                                                                                                                                                                                                                                                           RESULT 35
E01006/c
                                                                                                                                                                                                                                                                     VERSION
KEYWORDS
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                     COMMENT
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                                                                                                                         TITLE
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VERSION
KEYWORDS
SOURCE
ORGANISM
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Best Local Similarity
                                                                                                                                                                                                         Human herpesvirus 4 (Epstein-Barr virus)
Human herpesvirus 4
Viruses; dsDNA viruses, no RNA stage; Herpesviridae;
Gammaherpesvirinae; Lymphocryptovirus.
1 (bases 1 to 3400)
Hansu,Y.B.
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DNA sequences of the EBV genome, recombinant DNA molecules, processes for producing EBV-related antigens, diagnostic processes for producing EBV-related antigens, diagnostic grantanting sai
HANSU YOTSUTO BUORUFU
OS Epstein-Barr virus
PN JP 1986257188-A/2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 GGCTGGTGTCACCTGT 16
                                                                                                                  DNA SEQUENCE OF EBV GENOM, ANTIGEN AND DIAGNOSTIC COMPRODUCTION OF PREPARATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                           cDNA encoding p350 of Epstein-Barr virus. E01006 ·
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wolf, Hans Joachim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        synthetic construct synthetic construct
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                                                                                      Patent: JP 1986257188-A 2 14-NOV-1986;
                                                                                                                                                                                                                                                                                                                                                                                               E01006.1 GI:2169265
JP 1986257188-A/2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        compositions and pharmaceutical compositions containing said
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1 (bases 1 to 3400)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GGCTGGTGTCACCTGT 2664
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EP 0173254-A 3 05-MAR-1986;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /transI_table=11
/protein_id="CAA00938.1"
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DVNVTINFDVGGKKHQLDLDFGQLTPHTKAVYQPRGAFGGSENATNLFLLELLGAGEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SNLSMLVLQWASLAVLTLLLLLVMADCAFRRNLSTSHTYTTPPYDDAETYV" 1060 c 732 g 704 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    KPGEVNVTKGTPPQNATSPQAPSGQKTAVPTVTSTGGKANSTTGGKHTTGHGARTSTE
PTTDYGGDSTTPRPRYNATTYLPPSTSSKLRPRWTFTSPPVTTAQATVPVPPTSQPRF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                KWDNCNSTNITAVVRAQGLDVTLPLSLPTSAQDSNFSVKTEMLGNEIDIECIMEDGEI
SQVLPGDNKFNITCSGYESHVPSGGILTSTSPVATPIPGTGYAYSLRLTPRPVSRFLG
NNSILYVFYSGNGPKASGGDYCIQSNIVFSDEIPASQDWPTNTTDITYVGDNATYSVP
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encoding gp 250/350."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ALTMRSKKLPINVTTGEEQQVSLESVDVYFQDVFGTMWCHHAEMQNPVYLIPETVPYI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  organism="synthetic construct"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ocation/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 16;
Pred. No.
                                                                                                                                                      VOM, RECOMBINED DNA MOLECULE AND EBV RELATED COMPOSITION CONTAINING SAID ANTIGEN AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3400 bp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 3400;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PAT 29-SEP-1997
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BASE COUNT
ORIGIN
Search completed: August 15, 2003, 09:33:48 Job time: 503.975 secs
                                                                                                                                                 FEATURES
source
                                                                Query Match 88.9%; Score 16; DB 6; Length 3400; Best Local Similarity 100.0%; Pred. No. 14; Matches 16; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                 1 GGCTGGTGTCACCTGT 16
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2679 GGCTGGTGTCACCTGT 2664
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                                                                                                                904 a
                                                                                                                                                        3280. .3400 apolyA_site 3284. .3291. Location/Qualifiers
                                                                                                             .3400
                                                                     0;
                                                                   Gaps
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В

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Minimum DB seq length: 0
Maximum DB seq length: 200000000
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Post-processing: Listing first 120 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Word size :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Run on:
Pred. No. is the number of results predicted by chance to have a
                                                                                                                                                                                GenBmbl:*

1: gb ba:*

2: gb ht:*

3: gb in:*

4: gb ow:*

5: gb pat:*

7: gb pat:*

9: gb pr:*

9: gb pr:*

11: gb sts:*

11: gb sts:*

12: gb sy:*

13: gb un:*

6: em fun:*

6: em fun:*

7: em hum:*

8: em in:*

9: em ow:*

1: em or:*

1: em or:*

1: em ro:*

1: em ro:*

1: em ro:*

1: em ro:*

1: em ht:*

1: em ht:*

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1: em ht:*

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Kieff,E., Tanner,J., Hummel,M. and Beisel,C.
Vaccine against Epstein-Barr Virus
Patent: US 4707358-A 3 17-NOV-1987;
The University of Chicago; Chicago, IL
Location/Qualifiers
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Sorge, J.A. and Mullinax,R.L.
Polymerase compositions and uses thereof
Patent: US 555772-A 29 17-SEP-1996;
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Human herpesvirus 4
Viruses; dsDNA viruses, no RNA stage; Herpesviridae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Submitted (04-JUL-1996) S.D.V. Pepper, Paterson Institute for Cancer Research, Molecular Biology, Wilmslow Rd, Withington,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2 (bases 1 to 2663)
Pepper, S.D.V.
Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Immunisation of common marmosets with vaccinia virus expressing Epstein-Barr virus (EBV) gp340 and challenge with EBV J. Med. Virol. 50 (3), 263-271 (1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Epstein-Barr virus BLLF1 gene X99106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mackett, M., Cox, C., Pepper, S.D., Lees, J.F., Naylor, B.A., Wedderburn, N. and Arrand, J.R.
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  CCTTAGGAGGAACAAGTCCC 1844
                         CCTTAGGAGGAACAAGTCCC 20
                                                                               Conservative
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1 829 C
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GNSSTSTKPGEVNVTKGTPPKNATSPQAPSGQKTAVPTVTSTGGKANSTTGGKHTTGH
GARTSTEPTTDYGDDSTTPRPKYNATTYLPPSTSSKLRPRWTFTSPPVTTAQATVPVP
                                                                                                                                                                                                                                                                                                                                         VSGLGTAPKTILI TRTATNATTTTHKVI FSKAPESTTTSPTLNTTGFADPNTTTGLPS
STHVPTNLTAPASTGPTVSTADVTSPTPAGTTSGASPVTPSPSPRDNGTESKAPDMTS
PTSAVTTPTPNGTSPTPAMTTPTPNATSPTLGKTSPTSAVTTPTPNATSPTPAVTTPT
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SQVLPGDNKFNITCSGYESHVPSGGILITSTS PVATFIPGTAYSLLLTPRPVSRFLG
NNSILYVFYSGNGPKASGGDYCIQSNIVFSDEIPASQDMPTNTTDITYVGDNATYSPL
MVTSEDANSPNVTVTAFWAWPNNTETDFKCKWTLTSGTPSGCENISGAFASNRTFDIT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /translation="meaallvcoytioslihltgedpgffnveipefffyftcnvcta
DVNVTINFDVGGKKHQLDLDFGQLTPHTKAVYQPRGAFGGSENATNLFLLELLGAGEL
ALTWRSKKLPINVTTGEEQQVSLESUDVYFQDVFGTMKCHHAEMQNPVYLIPETVPYI
                                                                                                                                                                                                                                         PTSQPRFSNLSMLVLQWASLAVLTLLLLLVMADCAFRRNLSTSHTYTTPPYDDAETYV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /product="gp340"
/protein_id="CAA67558.1"
/db_xref="GI:2769560"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /db_xref="SPTREMBL:056854"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene="BLLF1"
codon_start=1
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strain="M81"
                                                                                             100.0%;
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Pred.
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                                                                               Indels
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E00513
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All128
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Matches 20
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Human herpesvirus 4
Viruses; dsDNA viruses, no RNA stage; Herpesviridae;
                                                                                                                                                                                                               3400 bp DNA linear PAT 07-E
Synthetic nucleotide sequence of the leftward reading frame
Bam L-fragment encoding gp 250/350.
All128
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Patent: JP 1985232094-A 1 18-NOV-1985;
UNIV CHICAGO:THE
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Genomic DNA encoding Epstein-Barr virus gp 220/200.
                                                        DNA sequences of the EBV genome, recombinant DNA molecules, processes for producing EBV-related antigens, diagnostic compositions and pharmaceutical compositions containing said
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 (bases 1 to 2721)
Eriotsuto, K., Jieroo
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              Patent: EP 0173254-A 3 05-MAR-1986; Wolf, Hans Joachim
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                                                                                                        Wolf,H.J
                                                                                                                                                    synthetic construct
                                                                                                                                                               synthetic construct
                                                                                                                                                                                               A11128.1 GI:490970
                                                                                                                     artificial sequences.
1 (bases 1 to 3400)
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hypothetical: No;
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30-JAN-1984 US 84 5753
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /product='gp 220/200'.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Human herpesvirus 4"
/mol type="genomic DNA"
/db xref="taxon:10376"
876 c 557 g 526 t
Location/Qualifiers
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KURISUTOFUAA
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 Human herpesvirus 4
Viruses; dsDNA viruses, no RNA stage; Herpesviridae;
Gammaherpesvirinae; Lymphocryptovirus.
1 (bases 1 to 3400)
                                                                                                                                                                                                                                                                                                                    DNA SEQUENCE OF EBV GENOM, RECOMBINED DNA MOLECULE ANTIGEN AND DIAGNOSTIC COMPOSITION CONTAINING SAID PRODUCTION OF PREPARATION
PALENC: JP 1986257188-A 2 14-NOV-1986;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human herpesvirus 4 (Epstein-Barr virus)
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JP 1986257188-A/2.
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23-AUG-1985 JP 1985185661
23-AUG-1984 EP 84 84110089, 23-AUG-1984 EP
U YOTSUTO BUORUFU
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Conservative 0; Mismatches
                                                                      topology: Linear; hypothetical: No;
                                                                                                                                                  C12N15/00,A61K39/245,C07H21/04,C07K13/00,C12N1/00,G01N33/569,G01N33/577//
                                                                                                                                                                                                                                                             Epstein-Barr virus
JP 1986257188-A/2
                                                      antı-sense: No;
                                                                                                             strandedness: Double;
                                                                                                                              C12P21/02, (C12N1/00, C12R1:19);
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a 1060 c 732 g 704 t
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ATNHTLGGTSPTPVVTSQPKNATSAVTTGQHNITSSSTSSMSLRPSSNPETLSPSTSD
NSTSHMPLLTSAHPTGGENITQVTPASISTHHVSTSSPAPRPGTTSQASGPGNSSTST
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VSGLGTAPKTLIITRTATNATTTTHKVIFSKAPESTTTSPTLNTTGFADPNTTTGLPS
STHVPTNLTAPASTGPTVSTADVTSPTPAGTTSGASPVTPSPSPWDNGTESKAPDMTS
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SQVLPGDNKFNITCSGYESHVPSGGILTSTSPVATPIPGTGYAYSLRLTPRPVSRPLG
NNSILYVFYSGNGPKASGGDYCIQSNIVFSDEIPASQDMPTNTTDITYVGDNATYSVP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /translation="meaallvcqytiqslihltgedpgffnveipefffyftcnvcta
DVNVTINfdvggkkhqldldfgqltphtkavyqprgafggsenatnlfllellgagel
ALIMRSKKLfINVTTGEEQQVSLESUDVYfqDVfgTmwchhaemqnpvylipetvfyi
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encoding gp 250/350."
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/db_xref="GI:490971"
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transl_table=
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/db_xref="taxon:32630"
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                                   Location/Qualifiers
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                                FEATURES
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Synthetic:
pURLP1.9.
A11178
                                         Patent:
Wolf, H
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Spaete, R. and Jackman, W.T.
Non-splicing variants of gp350/220
Patent: US 5824508-A 18 20-OCT-1998;
Location/Qualifiers
                                                                       DNA sequences of the EBV genome, recombinant DNA molecules, processes for producing EBV-related antigens, diagnostic compositions and pharmaceutical compositions containing sai
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  33
                                                                                                               artificial sequences
1 (bases 1 to 5019)
                                                                                                                                     synthetic construct synthetic construct
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1165 c 846 g
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/mol_type="genomic DNA"
/db_xref="taxon:10376"
1060 c 732 g 704 t
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/organism="synthetic construct"
/mol_type="genomic DNA"
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Pred. No. 1.
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Pred. No.
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5019 bp DNA IIIIEGT FAL 27-05 A. DNA sequence of hybrid peptide which contains beta-gelactosidase and p350 of Epstein-Barr virus.
                                                                                                                                                                                       DNA SEQUENCE OF EBV GENOM, RECOMBINED DNA MOLECULE AND EBV RELATED ANTICEN AND DIACHOSOSTIC COMPOSITION CONTAINING SAID ANTIGEN AND PRODUCTION OF PREPARATION PALENCE; JP 1986257188-A 3 14-NOV-1986;
                                                                                                                                                                                                                                                                                                                                                                               synthetic construct
synthetic construct
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1 (bases 1 to 5019)
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Artificial sequence;
14-NOV-1986
23-AUG-1985 JP 1985185661
23-AUG-1984 EP 84 84110089, 23-AUG-1984 EP 84 84110090 PI
                                                                                       JP 1986257188-A/3
                                                                                                                                                                    YOTSUTO BUORUFU
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NHPLMYTLCDRYGLYVVDEANIETHGMVPMNRLTDDPRWLPAMSERVTRMVQRDRNHP
SVIIWSLGNESGHGANHDALYRWIKSVDPSRPVQYEGGGADTTATDIICPMYARVDED
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SQPKNATSAVTTGQHNITSSSTSSMSLRPSSNPETLSPSTSDNSTSHMPLLTSAHPTG
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ATNATTTTHKVIFSKAPESTTTSPTLNTTGFADPNTTTGLPSSTHVPTNLTAPASTGP
TVSTADVTSPTPAGTTSGASPVTPSPSPWDNGTESKAPDWTSSTSPVTTPTPNATSPT
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YESHVPSGGILTSTSPVATPIPGTGYAYSLRLTPRPVSRFLGNNSILYVFYSGNGPKA
SGGDYCIQSNIVFSDEIPASQDMPTNTTDITYVGDNATYSVPMVTSEDANSPNVTVTA
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RIDPNAWVERWKAAGHYQAEAALLQCTADTLADAVLITTAHAWQHQGKTLFISRKTYR
IDGSGQMAITVDVEVASDTPHPARIGLNCQLAQVAERVNWLGLGPQENYPDRLTAACF
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rpsqqlrslngewrpawfpapeavpeswlecdlpeadtvvvpsnwqmjgydapiytnv
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PC C12N15/00,A61X39/2
PC G01N3/577//
PC C12P21/02,(C12N1/0
CC strandedness: Doub
CC topology: Linear;
CC hypothetical: No;
CC anti-sense: No;
CC *source: clone=pUH
FH Key LG
FT CDS
FT CDS
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Spacte, R. and Jackman, W.T.
Spacte, and Jackman, W.T.
Non-splicing variants of gp350/220
Patent: US 6458364-A 1 01-OCT-2002;
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AR233080
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1788 c 1433 g
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/mol_type="genomic DNA"
/mbl xref="taxon:32630"
/db xref="taxon:32630"
1529 c 1266 g 994 t
                                                                                                                                                                                                            Location/Qualifiers
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from patent US 6458364.
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/note='pBR322'.
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/note='poly cloning region'
3089. .4985
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85211017
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Human herpesvirus 4 (Epstein-Barr virus)
Human herpesvirus 4
Viruses; dsDNA viruses, no RNA stage; Herpesviridae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Two major outer envelope glycoproteins of Epstein-Barr virus are encoded by the same gene
J. Virol. 54 (3), 665-674 (1985)
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Kieff.E.
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ATNHTIGGTSPTPVVTSQDKNATSAVTTQHNITSSTSAVTTPIPNATGFTVGETSPQAN
ATNHTIGGTSPTPVVTSQDKNATSAVTTQHNITSSTSPERFRUTTSQASDETILSPSTSD
NSTSHMPLLTBSAHPTGGENITOVTPASISTHHVGTSSPERFRUTTSQASDGNSSTST
KPGEVNVTKGTPPQNATSPQAPSGQKTAVPTVTSTGGKANSTTTGGKHTTGHGARTSTE
PUTDYGGDSTTPRPKYNATTYLEPSTSSKLRPRWTFTSPVTTAQATVPVPPTSQPRF
SNLSHLVLQMASLAVLTLLLLLVMADCAFRRNLSTSHTYTTPPYDDAETYV"
join(1014 ...2514,3262 ...3737)
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SQVLPGDNKFNITCSGYESHVPSGGILITSTS PVANTPIPGTGAAVSLILTPRPVSRFLG
NNSILVVEYSGNGPKASGGDYCIQSNIVESDEIPASQDMPTNTTDITYGDNATYSVP
NTTSEDANS PNVTVTAFWAWPNNTETDFKCKWTLTSGTPSGCENISGAFASNRTFDIT
VSGLGTAPKTLIITRTANWATTTTHKVIFSKAPBSTTTSFTLMTTGFADPNTTTGLPS
STHVPTNLTAFASTGFTVSTAVVTSFTPAGTTSGASPVTPSBSBWDNCTESKAPDMTS
STHVPTNLTAFASTGFTVSTAVVTSTPTAGTSGASFVTPSBSBWDNCTSTSKAPDMTS
STSPVTTPTPAATSFTPAVTTFTFNATSFTPAVTTFTPNATSFTLGKTSFTSAVTTFT
                                             ATTYLPPSTSSKLRPRWTFTSPPVTTAQATVPVPPTSQPRFSNLSMLVLQWASLAVLT
LLLLVMADCAFRRNLSTSHTYTTPPYDDAETYV"
                                                                                           VSGLGTAPKTLI I TRTATNATTTTHKV I FSKAPESTTTSPTLNTTGFADPNTTTGLPS
STHVPTNLTAPASTGPTVSTADVTSPTPAGTTSGASPVTPSPSPWDNGTESTPPQNAT
SPQAPSGQKTAVPTVTSTGGKANSTTGGKHTTGHGARTSTEPTTDYGGDSTTPRPRYN
                                                                                                                                                           /translation="MEAALLYCQYTIQSLIHLTGEDPGPFNVEIPEFPPYPTCNVCTA
DVNVTINFDVGGKKHQLDLDFGGLTPHTAGYCPGAAFGGSENAFLFLELLGAGEL
ALTMRSKKLP INVTTGEEQQVSLESUDYYFQDVGGTMWCHHAEMQNPVLFLIFETVGAGEL
KMDNCNSTNITAVVRAQGLDVTLPLSLPTSAQDSNFSVKTEMLGNEIDIECIMEDGEI
SQVLFGDNKFNITGSGYESHVBSGGILTSTSPVATEIPGTGTGYASLRLTFRPVSRFLG
NNSLLYVFYSGNGFKASGGDYCIQSNIVFSDEIPASQDMFTYTDITYVGDNATYSPL
MYTSEDANSPNVTVTAFWAMPNNTETDFKCKWTLTSGTPSGCENISGAFASNRTFDIT
                                                                                                                                                                                                                                                                                                                       /codon_start=1
/protein_id="AAA45880.1"
/db_xref="GI:330361"
/note="major outer envelope glycoprotein gp220"
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dvnvtinfdvggkkhqldldfgqltphtkavyqprgafggsenatnlfllellgagel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /protein_id="AAA45881.1"
/db_xref="GI:330362"
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mol_type="genomic DNA"
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Human herpesvirus 4 complete
AJ507799
AJ507799.1 GI:23893576
  Jeang, K.T. and Hayward, S.D. Organization of the Epstein-Barr virus DNA molecule.
                                                                                                                Seguin, C., Farrell, P.J. and Barrell, B. DNA sequence and transcription of the B95-8 Epstein-Barr virus Mol. Biol. Med. 1 (3), 369-392 (1983)
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J. Cell Bianar
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Human herpesvirus 4
Viruses; dsDNA viruses, no RNA stage; Herpesviridae;
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                                                                                                                                                                                                                                                                                                                                                                                                                     Farrell, P.J., Deininger, P.L., Bankier, A. and Barrell, B. Homologous upstream sequences near Epstein-Barr virus proc. Natl. Acad. Sci. U.S.A. 80 (6), 1565-1569 (1983) 83169725
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Possible role of flanking nucleotides in recognition of the AUG initiator codon by eukaryotic ribosomes
Nucleic Acids Res. 9 (20), 5233-5262 (1981)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Griffin,B.E. Molecular cloning of the complete Epstein-Barr virus genome as set of overlapping restriction endonuclease fragments Nucleic Acids Res. 9 (13), 2999-3014 (1981)
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and Barrell, B.G.
DNA sequence analysis of the EcoRI Dhet fra
Epstein-Barr virus containing the terminal
Mol. Biol. Med. 1 (4), 425-445 (1983)
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Homology between two EBV early genes and HSV
reductase and 38K genes
Nucleic Acids Res. 12 (12), 5087-5099 (1984)
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A cis-acting element from the Epstein-Barr viral genome that permits stable replication of recombinant plasmids in latent infected cells
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The EB virus genome in Daudi Burkitt's lymphoma cells ha
deletion similar to that observed in a non-transforming
(P3HR-1) of the virus
EMBO J. 3 (4), 813-821 (1984)
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spliced Epstein-Barr virus gene expressed i
ymphocytes is created by circularization of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This sequence was assembled from B95-8 EBV [14] and Raji EBV [18] with sequence corrections [16, 19]. The number of major internal repeat units has been reduced from 11.6 [14] to a more typical 7.6 and the B95-8 deletion sequences have been restored to give a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (01-AUG-2002) Farrell P., Ludwig Institute for Cancer Research, Imperial College School of Medicine, St. Mary's Campu
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Virology 179 (1), 339-346 (1990)
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PRDDSSQHIYEBAGRGSMNPVCLPVIVAPYLFMLAA, TAASCFTASVSTVTATGLALS
LLLIAAVASSYAAAQRKLLTPVTUTTAVVTFFBAICLTMRIEDPPNSLLFALLAAAGE
LQGIYVLVMLVLLILAYRRRWRRLTVCGGIMFLACVLVLIVDAVLQLSPLLGAVTVVS
MTLLLLAFVLMLSSPGGLGTLGAALLTLAAALALLASLILGTLNLTTMFLLMLLMTLN
/codon_start=1
/product="terminal protein
/protein_id="CAD53383.1"
                                                                                                                                                                                                                VLLICSSCSSCPLSKILLARLFLYALALLLLASALIAGGSILQTNFKSLSSTEFIPNL
FCMLLLIVAGILFILAILTEWGSGNRTYGPVFMCLGGLLTMVAGAVWLTVMSNTLLSA
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/protein_id="CAD53382.1"
/db_xref="GI:23893577"
                                                                1280. .1495,1574.
/gene="LMP2"
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Jeang, K.T. and Hayward, S.D.
Organization of the Epstein-Barr virus DNA molecule. III. Lo.
Organization of the Epstein and characterization of the 1
of the P3HR-1 deletion junction and characterization of the 1
repeat units that form part of the template for an abundant 12-O-tetradecanoylphorbol-13-acetate-induced mRNA transcript
12-O-tetradecanoylphorbol-13-acetate-induced mRNA transcript
12-O-tetradecanoylphorbol-13-acetate-induced mRNA transcript
                                                                                                                                                                                                                                                                                                                                                                          4 (bases 1 to 172281)
Farrell, P.J., Deininger, P.L.,
Homologous upstream sequences
Proc. Natl. Acad. Sci. U.S.A.
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Arrand,J.R., Rymo,L., Walsh,J.E.,
Griffin,B.E.
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Bankier, A.T., Deininger, P.L., Farrell, P.J. and Barrell, B
Sequence analysis of the 17,166 base-pair EcoRI fragment
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Deininger, P.L., Bankier, A.,
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Transcription and DNA sequence of the
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Latent and lytic cycle promoters of
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                                                                                                                                   EMBO
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Nucleic Acids Res. 12 (
                          18 (bases 1 to 172281)
Farrell, P.J. and Barrell
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BamHI L fragment of B95-8
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This feature lists all occurences of the sequence AATAAA which is found normally approximately 20 bases upstream of the mRNA processing/polyA addition site. The rarely used homolog ATTAAA is only listed when it is found in a position close to the end of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        well as all the major open reading frames in the sequence. In general the term major is taken as the longest frame in a particular region taking into account the adjacent longest frames and likely transcription signals. Note that on this basis some long
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Direct Submission
Submitted (18-MAR-1988) Farrell P., Ludwig Institute for Cancer
Research, St. Mary's Hospital Medical School, Norfolk Place London
W2 1PG
W2 1PG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This is not a comprehensive listing of all such sequences and only the positions of a few have been noted because they occur in potentially interesting positions. The number quoted in the table is the position of the terminal base in the intron in each case.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              major reading frame.
SITEs of DONOR and ACCEPT sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The DNA sequence of B95-8 EBV has been revised [19]. The origina (Baer et al, 1984) base 359 has been deleted so the new sequence around that position reads TCAGTCTTT. To avoid renumbering the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Denotes the region that encompasses an origin of replication \mathsf{P}) \cdot \mathsf{113}] .
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P3HR1 and DAUDI with
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/db_xref="taxon:10376"
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DNA sequence and expression of the B95-8 Epstein-Barr virus genome Nature 310 (5974), 207-211 (1984)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Parker,B.D., Bankier,A., Satchwell,S., Barrell,B. and Farrell,P.J. Sequence and transcription of Raji Epstein-Barr virus DNA spanning the B95-8 deletion region Virology 179 (1), 339-346 (1990)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human herpesvirus 4
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Rickinson, A. and Kieff, E.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Original source text: Human herpesvirus 4 DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Restricted Epstein-Barr virus protein expression in Burkitt lymphoma is due to a different Epstein-Barr nuclear antigen transcriptional initiation site
Proc. Natl. Acad. Sci. U.S.A. 88 (14), 6343-6347 (1991)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human herpesvirus 4 (Epstein-Barr virus)
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Gammaherpesvirinae; Lymphocryptovirus.
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                                                                                                                                                                                                                                         /note="Overlap of B95-8 and deletion point (corresponds
                                                                                            /note="Overlap of B95-8 and Raji sequences at B95-8 deletion point (corresponds to 152,009-152,012 of V01555, and 11,832-11,835 of M35547)"
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                                                         /note="B95-8 sequences (corresponds to 152,013-172,282
                                                                                                                                                                                 note="Raji sequences (corresponds to 5-11,831 of M35547)"
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tifactual joining of B95-8 complete
Raji of the large deletion ביי
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to 152,009-152,012 in V01555,
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Unpublished
                                       Zhao, S., Dunn, D., von Niederhausern, A., Holt, R.A., Smith, H.O., Weinstock, G. and
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Rattus norvegicus (Norway rat)
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Direct Submission
Submitted (31-MAR-2003) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One maylor Plaza, Houston, TX 77030, USA
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Baylor Plaza, Houston,
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NOTE: This is a 'working draft' sequence. It currently consists of 49 contigs. The true order of the pieces is not known and their order in this sequence record is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence
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Chemistry: Dye-terminator Big Dye: 100% of reads
Assembly program: Phrap; version 0.990329
Consensus quality: 140865 bases at least Q40
Consensus quality: 147625 bases at least Q30
Consensus quality: 152515 bases at least Q20
Estimated insert size: 19404; sum-of-contigs estimation
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Center clone name: CH230-522E23
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Anderson, S., Arachchi, H.M., Barna, N., Bastien, V., Bloom, T.,

Boguslavkiy, L., Boukhgalter, B., Camarata, J., Chang, J., Choepel, Y.,

Collymore, A., Cook, A., Cooke, P., Corum, B., DeArellano, K.,

Piaz, J.S., Dodge, S., Dooley, K., Dorris, L., Erickson, J., Faro, S.,

Ferreira, P., FitzGerald, M., Gage, D., Galagan, J., Gardyna, S.,

Graham, L., Grand-Pierre, N., Hafez, N., Hagopian, D., Hagos, B.,

Hall, J., Horton, L., Hulme, W., Iliev, I., Johnson, R., Jones, C.,

Kamat, A., Karatas, A., Kells, C., Landers, T., Levine, R.,

Lindblad-Toh, K., Liu, G., Lui, A., Mabbitt, R., MacLean, C.,

Macdonald, P., Major, J., Manning, J., Matthews, C., McCarthy, M.,

Nguyen, C., Nicol, R., Norbu, C., O'Connor, T., O'Donnell, P.,

O'Neil, D., Oliver, J., Peterson, K., Phunkhang, P., Pierre, N.,

O'Neil, D., Oliver, J., Peterson, K., Phunkhang, P., Pierre, N.,

Roman, J., Schauer, S., Schupback, R., Seaman, S., Severy, P., Smith, C.,

Spencer, B., Stange-Thomann, N., Stojanovic, N., Stubbs, M.,

Talamas, J., Tesfaye, S., Theodore, J., Topham, K., Travers, M.,

Vassiliev, H., Venkataraman, V.S., Viel, R., Vo, A., Wilson, B., Wu, X.,

Vassiliev, H., Venkataraman, V.S., Viel, R., Vo, A., Wilson, B., Wu, X.,

Vannan D., Young, G., Zainoun, J., Zembek, L., Zimmer, A. and Zody, M.
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Direct Submission

All Submission

All
                                               Submitted (11-MAR-2003) Whitehead Institute/MIT Center for Genome Research, 320 Charles Street, Cambridge, MA 02141, USA On Mar 11, 2003 this sequence version replaced gi:28626751. All repeats were identified using RepeatMasker: All F.A. & Green, P. (1996-1997).
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http://ftp.genome.washington.edu/RM/RepeatMasker.html
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence as soon as it is available and the accession number will be preserved.
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67230
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequencing vector: Plasmid; n/a;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Center project name: L24242
Center clone name: 268_F_15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: sequence submissions@genome.wi.mit.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Center code: WIBR
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note="assembly_fragment"
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L9557. .39905
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193466: contig of 88597 bp in length
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Homo sapiens (human)
Homo sapiens
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human STS A005R48, sequence tagged site.
G20467
                                                                                                                      R24852,
                                                                                                                                                                                                                                                                                                                                  Buffer:
                                                                                                                                       Prepared with primer pairs derived from THC105423: GenBank Accession Numbers- H80345, H70110, F11453, F13120, R24326, R24340,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Protocol:
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Primer B: ATTGAGTGCTAGAGAATTACATA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (1996)
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Mark Adams
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Adams, M.D.
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67230. .104769
  /organism="Homo sapiens"
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                                                                                                                                                                                                                                                                                                                                                                      The Salk, Stanford, PGEC (SSP) Consortium members constructed and sequenced the pUNI (ORF) clones using the RAFL cDNAs: Yamada, K., Banh, J., Chan, M., Chang, C.H., Chang, E., Dale, J.M., Deng, J.M., Goldsmith, A.D., Lee, J.M., Onodera, C.S., Quach, H.L., Tang, C.C., Toriumi, M., Wu, H.C., Yamamura, Y., Yu, G., Bowser, L., Chen, H., Cheuk, R., Jones, T., Karlin-Neumann, G., Kim, C., Lam, B., Lin, J., Meyers, M.C., Miranda, M., Nguyen, M., Palm, C.J., Shinn, P., Southwick, A., Davis, R.W., Ecker, J.R. and Theologis, A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Direct Submission
Submitted (19-FEB-2002) Plant Gene Expression Center, 800 Buchanan
Street, Albany, CA 94710, USA
The RIKEN Genomic Sciences Center (GSC) members carried out the
collection and clustering of RAFL CDNA; (RAFL CDNA; "RIKEN
Arabidopsis Full-Length cDNA'): Seki,M., Narusaka,M., Ishida,J.
Satou,M., Kamiya,A., Sakurai,T., Carninci,P., Kawai, J.,
Hayashizaki,Y. and Shinozaki,K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Yamada, K., Banh, J., Chan, M.M., Chang, C.H., Chang, E., Dale, J.M., Deng, J.M., Goldsmith, A.D., Lee, J.M., Onodera, C.S., Quach, H.L., Tang, C.C., Toriumi, M., Wu, H.C., Yamamura, Y., Yu, G., Bowser, L., Carninci, P., Chen, H., Cheuk, R., Hayashizaki, Y., Ishida, J., Jones, T., Kamiya, A., Karlin-Neumann, G., Kawai, J., Kim, C., Lam, B., Lin, J., Meyers, M.C., Miranda, M., Narusaka, M., Nguyen, M., Palm, C.J. Sakurai, T., Satou, M., Seki, M., Shinn, P., Southwick, A., Shinn, P., Southwick, A., Baris, R.W., Ecker, J.R. and Theologis, A.
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Yamada, K., Liu, S.X., Sakano, H., Pham, P.K., Banh, J., Etgu, P., Yamada, K., Liu, S.X., Sakano, H., Pham, P.K., Banh, J., Etgu, P., Lee, J.M., Toriumi, M., Yu, G., Brooks, S., Chao, Q., Chen, H., Karlin-Neumann, G., Kim, C., Lam, B., Miranda, M., Nguyen, M., Rarlin-Neumann, G., Kim, C., Lam, B., Miranda, M., Nguyen, M., Rarlin-Neumann, G., Kim, C., Lam, B., Miranda, M., Nguyen, M., Rarlin-Neumann, G., Kim, C., Lam, B., Miranda, M., Nguyen, M., Rarlin-Neumann, G., Kim, C., Lam, B., Miranda, M., Nguyen, M., Rarlin-Neumann, G., Kim, C., Lam, B., Miranda, M., Nguyen, M., Rarlin-Neumann, G., Kim, C., Lam, B., Miranda, M., Nguyen, M., Nguyen,
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                                                                                                                                                                                                                                                                          Yamada, K. (SSP/PGEC) and Seki, M. (RIKEN GSC) contributed equally to this work. Shinozaki, K. (RIKEN GSC) and Theologis, A. (SSP
                                                                                                                                      Annotation is based on the January genome submitted to GenBank.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Arabidopsis Open Reading Frame (ORF) Clones
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                                                                                                      Location/Qualifiers
/organism="Arabidopsis thaliana"
/mol_type="mRNA"
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                                                                                                                                                             2 (bases 1 to 1767)

Yamada, K., Liu, S.X., Sakano, H., Pham, P.K., Banh, J., Chung, M.K., Dale, J.M., Gibson, H.A., Goldsmith, A.D., Jiang, P.X., Lee, J.M., Dale, J.M., Tang, C.C., Toriumi, M., Yu, G., Bowser, L., Carninci, P., Chen, H., Cheuk, R., Hayashizaki, Y., Ishida, J., Jones, T., Kamiya, A., Karlin. Neumann, G., Kawai, J., Kim, C., Koesema, E., Lam, B., Lin, J., Meyers, M.C., Mizanda, M., Narusaka, M., Nguyen, M., Palm, C.J., Sakurai, T., Satou, M., Seki, M., Shinn, P., Southwick, A., Tracy, S.E., Sahinozaki, K., Davis, R.W., Ecker, J.R. and Theologis, A.
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(bases 1 to 1767)
Yamada, K., Liu, S.X., Sakano, H., Pham, P.K., Banh, J., Chung, M.;
Goldsmith, A.D., Lee, J.M., Quach, H.L., Toriumi, M., Yu, G., Bow,
Garninci, P., Chen, H., Cheuk, R., Hayashizaki, Y., Ishida, J.,
Jones, T., Kamiya, A., Karlin-Neumann, G., Kawai, J., Kim, C., Lay
Jones, T., Kamiya, A., Karlin-Neumann, G., Kawai, J., Kim, C., Lay
Jones, T., Sakina, Narusaka, M., Nguyen, M., Palm, C.J., Sakur,
Satou, M., Seki, M., Shinn, P., Southwick, A., Shinozaki, K.,
Davis, R.W., Ecker, J.R., and Theologis, A.
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Street, Albany, CA 94710, USA RIKEN Genomic Sciences Center
                                         Submitted (07-JUN-2001) Plant Gene Expression Center, 800 Buchanan Street, Albany, CA 94710, USA
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Arabidopsis thaliana
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                                                                                                                                  Direct Submission
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/protein id="ALIS5111.1"
/protein id="ALIS5111.1"
/db_xref="Gi:19310761"
/translation="MSYLLRRNKFVALLKRQSLIFPITSTEAKTLINPDPNIPQFQNP
/translation="MSYLLRRNKFVALLKRQSLIFPITSTEAKTLINPDPNIPQFQNP
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    (GSC) members carried out the
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2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PLN 18-SEP-2002
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DEFINITION
ACCESSION
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AX151433
                              SOURCE
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Best Local Similarity
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                                                                                            AX151433
Sequence 38 from Patent
AX151433
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  collection and clustering of RAFL CDNAs (RAFL CDNA: 'RIKEN Arabidopsis Full-Length cDNA'): Seki,M., Narusaka,M., Ishida Satou,M., Kamiya,A., Sakurai,T., Carninci,P., Kawai,J., Hayashizaki,Y. and Shinozaki,K.
Shrimp white spot syndrome virus Shrimp white spot syndrome virus
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ecotype: Columbia"
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CSIFR AHYATQSSKF PEYEMPTVTWGVIQKKEKLVNRVKICDYLKGLGITTDELES
IELPSTIEVMCERVEFLAKLGLTIDDINEYPHALGCSVRKNLLFVLAKLEKIGISTSK
IGEFVKNYPQVLHASVVVELAFVVKFLRGLDVEKQDLGYVLMKYFELLGFKLEGTMST
SVAYLNSIGVSFRDIGPMVTQSVFYLLGMRVGTMIKFLVDYLISIGLFKKIVARMLEKR
SYIGYNLETTVKRNVDCLISFGVFKKELLFLLLAQYPQDLIGLFVKAKMSTQQYFFSLK
KSIUGYNLETTVKRNVDCLISFGVKKELLFLLLAQYPQDLIGLFVKAKMSTQQYFFSLK
LKIDPEGFARVVEKMPQIVSLKQNVIMKPIEFLLGRAFQVEDIAKMVVRCPQILCSRV
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/protein_id="AAK64024.1"
/db_xref="GI:14532592"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /gene="At2g44020"
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126. .1649
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RESULT 22
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PE Corporation (NY) (US); The Third Institute of Oceanography, State Oceanic Administration (CN); Sinogenomax Co., Ltd. (CN)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Xu,X., Ya
Kodira,C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Miyajima,N., Tanaka,A., Kotani,H., Nomura,N. and Ohara,O. Prediction of the coding sequences of unidentified human genes. VII. The complete sequences of 100 new cDNA clones from brain which can code for large proteins in vitro DNA Res. 4 (2), 141-150 (1997)
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                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (28-MAR-1997) Osamu Ohara, Kazusa DNA Res
1532-3, Yana, Kisarazu, Chiba 292-0812, Japan
(E-mail:Cdnainfo@kazusa.or.jp, Tel:+81-438-52-3913)
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human mRNA for KIAA0382
AB002380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Direct Submission
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Ohara,O., Nagase,T., Kikuno,R. and Nomura,N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens (human)
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/mol type="genomic DNA"
/db_xref="taxon:92652"
/db_7 c 526 g 596 t
/protein_id="Baa20836.1"
/db_xref="G1:2224705"
/db_xref="G1:2224705"
/tdb_xref="G1:2224705"
/tabslation="Leyterahvrtlkvldovfyorvsregilspselrkifsnledi
/tabslation="Leyterahvrtlkvldovfyorvsregilspselrkifsnopfale
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MIKSROKKDSRFOTFVODAESNPLCRRLOLKDIIFTOMORLTKXFLLLDNIAKYTEWP
TERRKVKKAADHCROILWFVNOAVKEABNKORLEDYORKDTSLKLGSFYPNVEELN
LDLTKRKMIHEGPLVWKVNRDKTIDLYTLLLEDILVLLOKODDRLVLRCHSKILASTA
                                                                                                                                                                                                                                                                                                                             /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                   /gene="KIAA0382"
                                                                                                                                                                                                        /gene="KIAA0382"
                                                                                                                                                                                                                                                clone_
                                                                                                                                                                                                                                                                                                              clone="HH0685"
                                                                                                                                            codon_start=3
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                                                                                                                                                                                                                                              tissue_type="brain"
clone_lib="pBluescriptII SK plus"
                                                                                                                                                                                                                                                                                      sex="male"
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Pred. No.
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LISSKPQSHSLSTSGKSEVRDLFVAERQFAKEQHTDGTLKEVGEDYQIAIPDSHLPVS EERWALDALRNIGLLKQLLVQQLGLTEKSVQEDWQHFPRYRTASQGPQTDSVIQNSEN IKAYHSGEGHMPFRTGTGDIATCYSPRTSTESFAPRDSVGLAPQDSQASNILVMDHMI

MTPEMPTMEPEGGLDDSGEHFFDAREAHSDENPSEGDGAVNKEEKDVNLRISGNYLIL

DSKHTFSPVIKLSTVLVRQVATDNKALFVISMSDNGAQIYELVAQTVSEKTVWQDLIC RMAASVKEQSTKPIPLPQSTPGEGDNDEEDPSKLKEEQHGISVTGLQSPDRDLGLEST

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1 (bases 1 to 9501)
Kourlas, P.J., Strout, M.P., Becknell, B., Veronese, M.L., Croce, C.M., Theil, K.S., Krahe, R., Ruutu, T., Knuutila, S., Bloomfield, C.D. and
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Homo sapiens guanine nucleotide exchange factor (LARG) mRNA,
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Kourlas, P.J., Strout
Caligiuri, M.A.
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QQRWGAMEYSCFEIQSPSSCADSQSQIMEYIHKIEADLEHLKKVEESYTILCQRLAGS
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                  RLQLLÕEDYNRTPAQRLLKEIQEAKKHIPQLQEQLSKATGSAQDGAVVTPSRPLGDTL
TVSEAETDPGDVLGRTDCSSGDASRPSSDNADSPKSGPKERIYLEENPEKSETIQDTD
                                                       DPTDSSSKKTKSSSEESRSEIYGLVQRCVIIQKDDNGFGLTVSGDNPVFVQSVKEDGA
AMRAGVQTGDRIIKVNGTLVTHSNHLEVVKLIKSGSYVALTVQGRPPGSPQIPLADSE
VEPSVJGHMSPIMTSPHSPGASGNMERITSPVLMGEENNVVHNQKVEILRKMLQKEQE
                                                                                                                                    /codon_start=1
/product="guanine nucleotide exchange factor"
/protein_id="AAF36817.1"
/db_xref="GI:7110160"
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TQSLVGSPSTRIAPHIIGAEDDDFGTEHEQINGQCSCFQSIELLKSRPAHLAVFLHHV
                                                                                                                                                                                                                 gene="LARG"
                                                                                                                                                                                                                                                                                                                                                                              organism="Homo sapiens"
|mol_type="mRNA"
                                                                                                                  translation="MSGTQSTITDRFPLKKPIRHGSILNRESPTDKKQKVERIASHDF
                                                                                                                                                                                                                                                      gene="LARG"
                                                                                                                                                                                                                                                                             tissue_type="prostate" . .9501
                                                                                                                                                                                                                                                                                                                  map="11q23.3"
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2 (bases 1 to 16529)

Hattori,M., Ishii,K., Toyoda,A., Taylor,T.D., Hong-Seog,P., Fujiyama,A., Yada,T., Totoki,Y., Watanabe,H. and Sakaki,Y. Direct Submission
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                                                                                                                                                                                                                                                                                                                                                        (E-mail:hattori@gsc.riken.go.jp, URL:ht
Tel:81-45-503-9111, Fax:81-45-503-9170)
On Jun 12, 2003 this sequence version r
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                                                                                                                                                                                                                                                                                                                                                                                                                                  and Chemical Research (RIKEN), Genomic Sciences Center (GSC);
1-7-22 Suehiro-chou, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Submitted (13-MAY-1999) Masahira Hattori, The Institute of Physical
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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ERAHWETLKVLDQVFYQRVSREGILSPSELRKIFSNLEDILQLHIGLMEQMKAVRKRN
ETSVIDQIGEDLLTWFSGFGEEKLKHAAATFCSNQPFALENEM
AESNPLCRRLQLKDIIFOMQRLTKYELLLDNIAKYTEWPTEREKVKKADSRFQTFVQD
AESNPLCRRLQLKDIIFOMQRLTKYELLLDNIAKYTEWPTEREKVKKADHCRQILM
YVNQAVKEAENKORLEDYQRRLDTSSLKLSBYPNVEELRNLDLTKRKMIHEGPLVWKV
MRDKTIDLYTLLLEDILVLLOKODDRLVLRCHSKILASTADSKHTFSSPVIKLSTVLVR
QVATDNKALFVISMSDNGAQIYELVAQTVSEKTVWQDLICRMAASVKEOSTKPIPLPQ
STPGEGONDEEDPSKLKEEGHGISVTGLOSDRDLGLESTILSSKPOSHSTKPIPLPQ
STPGEGONDEEDPSKLKEEGHGISVTGLOSDRDLGLESTILSSKPOSHSTKHIGHLKQL
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TLYQCLGLTEKSYQEDWOHFPKYRTASGGFQTDSVIQNSENIKAYHSGEGHMPFRTGTG
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RRPELLPEDLHRHY I QTMQERVHJEVQORLEDPRQKRSMGLTLAESELTKLDAERDKD
RLTLEKERTCAEQI VAKIEBULMTAQAVEEDKSSTMQYVILMYMKHLGVVKVEPRNLE
HKRGRI IGFLPKI KQSMKKDKEGEEKGKRRGFPS I LGPPRRPSRHDNSAI GRAMELQKA
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LTLQPMTGIPAVESTHQQQHSPQNTHSDGAISPFTPEFLVQQRWGAMEYSCFEIQSPS
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
/chromosome="21"
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                                                                                                                         /clone="CMF21-74A4"
3739 c 3784 g
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ne 21 clone:CMF21-74A4,
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<pre>repear_regrou / rpt_family="Alu" /rpt_type=dispersed  variation</pre>	ion 6	<pre>variation 612 /frequency="0.36" /replace="q"</pre>	/replace="t" 583 /frequency="0.18" /replace="t"	/ipt_tamily="main" /rpt_type=dispersed variation 523 /frequency="0.01"	/freq /repl	/frequency="0.05" /replace="c"	variation 322 ""  /frequency="0.18" /replace="c" variation 353	variation 297  /frequency="0.05"  /frequency="0.05"	variation // //reqlency="0.03" //replace="a"	12.	/or /db/	(URL: http://egp.gs.washington.edu). FEATURES Location/Qualifiers source 132813		e Scie	AUTHORS Rieder, M.J., Livingston, R.J., Braun, A.C., Montoya, M.A., Chung, MW., Miyamoto, K.E., Nguyen, C.P., Nguyen, D.A., Poel, C.L., Robertson, P.D., Schackwitz, W.S., Sherwood, J.K., Witrak, L.A. and Nickerson, D.A.	Eukaı Mamma	m m	ON Homo sapiens X-ray repair complementing defective re hamster cells 2 (XRCC2) gene, complete cds. N AF520762 1 GI:21489901	RESULT 25 AF520762 AF520762 AF520762 AF520762 AF520762 AF520762 AF520762	5368 AGGAGGAACAAGTCCC	Qy 5 AGGAGGAACAAGTCCC 20
/replace="c" /replace="c" /replace="xRCC2" /frequency="0.02" /replace="a"	/replace="g" variation 2150 /gene="xCC2"	variation 2002 /gene="XRCC2" /frequency="0.03"	<pre>variation 1934 /gene="XRCC2" /frequency="0.05" /replace="c"</pre>	/gene="XRCC2" /frequency="0.02" /replace="a"		)	/translation="MCSAFHRAESGTELLARLEGRSSLKEIEPNLFADEDSPVHGDIL EFHGPEGGKTEMLYHLTARCILFKSEGGLEVEVLFIDTDYHFDMLRLVTILEHRLSQ SSEEIKKYLLAFFLYVCSSGTHLLLTLYSLESBHCSLEHLDDLSAFFWIDRV NGGESVNLQESTLRKCSQCLEKLVNDYRLVLFATTQTIMQKASSSSEEPSHASRRLCD	,	<pre>/gene="%xCC2" /codon_starts repair complementing defective repair in /product="%x-ray repair complementing defective repair in</pre>	/replaces / 1624,1688516966,2834029061)	Chinese hamster cells 2" variation 1517 /gene="XRCC2" /from="XRCC2"	)01624,16885 (RCC2" :="X-ray repair	1500. 31217 /gene="XRCC2"	variation 1498  /frequency="0.18"	/replace="t" variation 1389 /frequency="0.01" /replace="c"	/replace="t" variation 1334 /frequency="0.04"	/replace="" 171 /frequency="0.01"	misc_feature 1030. 1130 /note="Region not scanned for variation" variation 1133 .1136 /frequency="0.91"	/replace="g" variation 977 /frequency="0.05" /replace="c"	/replace="a" variation 964 /frequency="0.05"	variation 924 /frequency="0.04"

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RESULT 26
                                                                                                                       Matches
                                                                                                                                       Query Match
Best Local
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                                                     24451 AGGAGGAACAAGTCCC 24466
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                                                                                     5 AGGAGGAACAAGTCCC 20
                                                                                                                                         Similarity
                                                                                                                       Conservative
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/rpt_type=dispersed
3047. .3331
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201. .4476
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replace="c"
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                                                                                                                    80.0%; Score 16; DB 9; Length 32813; 100.0%; Pred. No. 1e+02; Tindels
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VERSION
KEYWORDS
SOURCE
ORGANISM
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AUTHORS
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Unpublished

3 (Dases 1 to 41308)

Ricke,D.O., Bruce,D., Mundt,M., Doggett,N., Munk,C., Saunders,E.,
Robinson,D., Jones,M., Buckingham,J., Chasteen,L., Thompson,S.,
Goodwin,L., Bryant,J., Tesmer,J., Meincke,L., Longmire,J.,
White,S., Ueng,S., Tatum,O., Campbell,C., Fawcett,J., Maltbie,M.,
White,S., Land Deaven,L.
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AC005203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (30-JUN-1998) Center for Human Genome Institute, Los Alamos National Lab Alamos, NM 87545, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 (bases 1 to 41308)
Ricke,D.O., Bruce,D., Mundt,M., Doggett,N., Munk,C., Saunders,E.,
Robinson,D., Jones,M., Buckingham,J., Chasteen,L., Thompson,S.,
Goodwin,L., Bryant,J., Tesmer,J., Meincke,L., Longmire,J.,
White,S., Ueng,S., Tatum,O., Campbell,C., Fawcett,J., Maltbie,M.,
Misra,M. and Deaven,L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 41308)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Large Scale Sequence Analysis and Annotation with the Sequence Comparison Analysis (SCAN) System
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ricke, D.O.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AC005203.1 GI:3273381
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Homo sapiens chromosome 16, cosmid clone RT70 (LANL), complete
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens (human)
                                                                                                                                    complement (1433...1698)
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                                                                                                                                                                                                                                                                                                                            /rpt_family="Alu" 654. .686
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   complement(4. .321)
/note="97% identity EST
/db_xref="dbEST:W70228"
                                                                                                                                                                                                                                                          /rpt_unit=A
717. .981
                                                                                                                                                                                                                                                                                                                                                           /db_xref="dbSTS:G37491"
44. .213
                                                                  /note="DNaseI"
<1747. .1889
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  complement (4. .321)
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                                                                                                    .747. .4885
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      clone="RT70"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /chromosome="16"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           location/Qualifiers
                                                   'gene="DNaseI"
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Laboratory, MS M888, Los
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/note="88% & 83% identity mouse dbEST:AA060574"
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26621. .26692,32898. .33038,35957. .36039,37729. .>37887)
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complement(5177. .5342)
                                                                                                                                                                                                                                                                                                                                                           complement (join (8887. .9035, 9772. .9863))
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5347. .5600
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DAVAEIDALYDVYLDVQEKWGLEDVMLMGDFNAGCSYVRPSQMSSIRLWTSPTFQWLI
PDSADTTATFTHCAYDRIVVAGMLLRGAVVFDSALPFNFQAAYGLSDQLAQAISDHYP
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7643. .7846
                                                                                                                                                                                                                                                                                                                                                                                                                                         /rpt_unit=TAGA
7223. .7360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     complement (6633.
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/gene="TRAP1"
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4939. .5135
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3539. .3654
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                                                                                                                                                 protein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /gene="DNaseI"
1900. .4012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /product="DNaseI"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            genes"DNaseI
                                                                                                                                                                     note="tumor necrosis factor type 1 receptor associated"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene="TRAP1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   'gene="TRAP1"
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'gene="DNaseI"
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                                                                                                                                                                                                                                                                                                                                       gene="TRAP1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  rpt_type=tandem
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        'note=" (TAGA) 8"
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db_xref="dbSTS:G06033"
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AC087682 AC087682.1 GI:12229457 HTG; HTGS\_PHASE0.

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RESULT 27
AC087682/c
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Homo sapiens chromosome 18 clone RP11-886K22 map 18, LOW-PASS SEQUENCE SAMPLING.
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                                                                                                                                                                                            Similarity
                                                                                                                                  AGGAGGAACAAGTCCC 20
                                                                                                             AGGAGGAACAAGTCCC 34191
                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                     /rpt_family="Alu"
complement(19602.
/gene="TRAP1"
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/gene="TRAP1"
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AEASSKIIGQFGVGFYSAFMVADRVEVYSRSAAPGSLGYQWLSDGSGVFEIAEASGVR
TGTKIIIHLKSDCKEESSEAAVRDVVTKYSNEVSFPLYLNGRRMNTLQAIMMDPKDV
REWQHEEFYRYVAQAHDKPRYTLHYKTDAFLINIRSIFYYPDGRRMFTLAIMFELGSSVA
LYSRKVLIGTKATDILFWHJRFIRGVVDSEDIPLNLSRELLOBSALIRKLRUVLQQRL
IKFFIDQSKKDAEKYAKFFEDYGLFMREGIVTATEQEVKEDIAKLLRYESSALPSGQL
TSLSEYASBWRAGTENIYYLCAPNHHLAEHSPYYEAMKKKDTEVLFCTEQFDELTTLH
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/note="87% & 80% identity
complement (9783. .9868)
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VKVTLRLDTHPAMVTVLEMGAARHFLRMQQLAKTQEERAQLLQPTLEINP"
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18932. .19247
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complement(18624.
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Pred. No. 98;
                                dq 68589
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·	COMMENT	TITLE	SOURCE ORGANISM REFERENCE AUTHORS TITLE JOURNAL REFERENCE AUTHORS
1 714: contig of 714 bp in length 1 814: gap of 100 bp 1 1522: contig of 708 bp in length 1 1523: l622: gap of 100 bp 1 1623: 2330: contig of 708 bp in length 2 2331: 2430: gap of 100 bp 2 4331: 3132: contig of 702 bp in length 3 133: 3232: gap of 100 bp 3 133: 3232: gap of 100 bp 3 133: 3944: contig of 712 bp in length 4 4045: 4044: gap of 100 bp 4 4045: 4741: contig of 697 bp in length 4 4841: gap of 100 bp 5 570: contig of 729 bp in length 5 571: 5670: gap of 100 bp 5 6481: gap of 100 bp 6 6488: 7189: contig of 702 bp in length		H HUTS. MOH. IOHS MIM S	Vertebrata, E i; Hominidae, l nder,E. 886K22 r,E., Allen,N.
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RESULT 28
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                             AUTHORS
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Best Local :
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                                                                                                                                                                 Human DNA from overlapping chromosome 7 PAC and Pl clones
             Brookman, K.W.,
                                                       Eukaryota;
Mammalia; I
                                                                                                                                        containing
AC003109
                             Liu,
                                                                                  Homo sapiens
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                                      karyota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
mmalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
(bases 1 to 76850)
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           N., Lamerdin, J.E., Tebbs, R.S., Schild, Dskman, K.W., Siciliano, M.J., Walter, C.A.,
                                                                                               sapiens (human)
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Submitted (17-MAR-1998) Human Genome Center, Lawrence Livermore National Laboratory, 7000 East Ave., Livermore, CA 9451, USA Sequence derived from PAC clone 13620 from bases 1 to 10,382, and from P1 clone 7515 from bases 10,095 to 76,850. No polymorphisms discrepancies were detected in the overlapping 288 base pairs.
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Submitted (19-NOV-1997) Human Genome Center, Lawrence Livermore
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Thompson,L.H.
XRCC3, New Human Rad51- Family Members, Promote Chromosome Stability and Protect Against DNA Crosslinks and Other
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bases 1 to 76850)
                                                                         /product="Xrcc2"
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/protein_id="AAC05802.1"
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SEFEIIKYCLGRFFLLYCSSSTHLLLTLYSLSSEFCSHPSLCLLLILDSLSAFYWIDRV
NGGESYNLOSSTLRKCSQCLEKLVNUYRLVLFATTQTIMQKASSSSEEPSHASRRLCD
VDIDYRPYLCKAMQQLVKHRMFFSKQDDSQSSNQFSLVSRCLKSNSLKKHFFIIGESG
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complement(3549.
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/rpt_family="HSATI"
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                          gene="XRCC2"
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complement(16576..16)
/rpt_family="Alu"
17529..17829
   /rpt_family="Alu"
28597. .28774
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complement(6510...
/rpt_family="Alu"
co/complement(family="Alu")
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/rpt_family="Alu"
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21931. .22256
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/rpt_family="Alu"
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/note="BLASTX similarity to
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complement(10534. .10574)
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Patent: EP 1270724-A 607 02-JAN-2003;
National Institute of Advanced Industrial Science and Te
                                                                                                                                          Suwa, M., Asai, K., Akiyama, Y. and Aburatani, H
                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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complement(28851.
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29170. .29452
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607 from Patent EP1270724.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="BLASTN similarity to Y08837 (1. .336); match: 0.99, score: 7.0e-130; database searched: nt; H.sagiens mRNA for RAD51-like protein-Other overlapping matches: BLASTX similarity to PID (93. .149, 206. .243); match: 0.43, score: 9.7e-06; database searched: nr; (Y13144) Rad51 homologue [Trypanosoma bruce1]-BLASTX similarity to 2262211 (101. .162); match: 0.43, score: 1.8e-06; database searched: nr; (U92068) RecA-like protein [Mus musculus] "complement (31829. .32256)
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31037. 3107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note="DDS similarity to AA255626 zs31f06.sl NCI_CGAP_GCB1 Homo sapiens CDNA clone 686619 3' similar to contains Alu repetitive element;contains element TR6 repetitive element;(186..427); 96% identity." Complement (186396..36646) /rpt_family="Alu" 36742..37029
/organism="Homo sapiens"
/mol_type="genomic DNA"
                                             Location/Qualifiers
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36278. .36522
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33635. .33888
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d, score: 67.000"
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Best Local Similarity
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                                                                                                                                                   finding wemensurver.

transmembrane helix prediction.

And the sequence is submitted by the collaborative project between [Computational Biology Research Center (CBRC), National Institute of Advanced Industrial Science and Technology (AIST)] and [Genome Science Division, Research Center for Advanced Science and Technology (ROST), University of Tokyo].
                                                                                                                                                                                                                                                                                                Submitted (11-JUL-2001) Makiko Suwa, Computational Biology Research Center (CBRC), National Institute of Advanced Industrial Science and Technology (AIST); 2-41-6 Ammi Koto-ku, Tokyo 135-0064, Japan (E-mail:m-suwa@aist.go.jp, UKL:http://www.cbrc.jp/, Tel:81-3-3599-8080, Fax:81-3-3599-8081)
Tel:81-3-3599-8080, Fax:81-3-3599-8081)
This sequence is a seven transmembrane helix receptor candidate predicted from the whole human genome sequences using our automated system that contains programs of gene finding (GeneDecoder), sequence search, motif-domain assignment and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
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58541. .58638,60172. .60304,62427. .62580,71080. .71140.
72348. .72417,75647. .75738,77542. .77668,82828. .82869
84170. .84310)
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RHFLEDVGCWDINANASIWWIIRGPVILSILINFILFINILRILMRKLRTQETRGNEV
SHYKRLARSTLLLIPLFGIHYIVFAFSPEDAMEIQLFFELALGSFQGLVVAVLYCFLN
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clqelsreqtgdlgteqpvpgcegmwnuscereketsvllkckymytvgyssslymtlvs
lfrncttqdgwsettprrhlacgvnyndssereketsyllkckymytvgyssslymtlvs
lgilcafrrlhctrnyihmhlfvsfilralsnfikdavlfssddvtycdahragcklv
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/protein_id="CAD69336.1"
/db_xref="GI:28798797"
                      /db_xref="taxon:9606"
/chromosome="2"
                                                                mol_type="genomic DNA"
isolate="CBRC7TM_223"
                                                                                                           organism="Homo sapiens"
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100.0%; Pred. No.
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CE 2 (bases 1 to 94357)

Birren, B., Linton, L., Nusbaum, C., Lander, E., Allen, N., Anderson, M., RS Birren, B., Linton, L., Beckerly, R., Boguslavkiy, L., Boukhgalter, B., Baldwin, J., Barna, N., Beckerly, R., Boguslavkiy, L., Boukhgalter, B., Brown, A., Castle, A., Colangelo, M., Collins, S., Collymore, A., Cooke, P., Deartellano, K., Dewar, K., Domino, M., Donelan, L., Doyle, M., Ferreira, P., FitzHugh, W., Forrest, C., Funke, R., Gage, D., Galagan, J., Gardyna, S., Grant, G., Hagos, B., Heaford, A., Horton, L., Howland, J.C., Johnson, R., Jones, C., Kann, L., Karatas, A., Klein, J., Hewland, J.C., Johnson, R., Jones, C., Kann, L., Karatas, A., Klein, J., McEwan, P., McGurk, A., McKernan, K., McLaughlin, J., Marquis, N., McEwan, P., McGurk, A., McKernan, K., McLaughlin, J., Meldrim, J., Peterson, K., Pollara, V., Riley, R., Roy, A., Santos, R., Severy, P., Stange-Thomann, N., Stojanovic, N., Subramanian, A., Talamas, J., Tesfaye, S., Tirrell, A., Vassiliev, H., Vo, A., Wheeler, J., Wu, X., Nicosion, S., McKernan, A., Talamas, J., Talamas, J., Zimmer, A., and Zody, M.
                                                                                                                                                                                                                                                                                                                                                        Submitted (28-SEP-1999) Whitehead Institute/MIT Center for Genome Research, 320 Charles Street, Cambridge, MA 02141, USA ON OCt 1, 1999 this sequence version replaced gi:5931399.
All repeats were identified using RepeatMasKer: Smit, A.F.A. & Green, P. (1996-1997)
http://ftp.genome.washington.edu/RM/RepeatMasker.html.
*NOTE: This record contains 98 individual
*sequencing reads that have not been assembled into
*contigs. Runs of N are used to separate the reads
*and the order in which they appear is completely
*arbitrary. Low-pass sequence sampling is useful for
*identifying clones that may be generich and allows
*overlap relationships among clones to be deduced.
*However, it should not be assumed that this clone
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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94357 bp DNA linear HTV
Homo sapiens clone 11_J_20, LOW-PASS SEQUENCE SAMPLING
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RHFLEDVGCWDINANASIWWIIRGPVILSILINFILFINILRILMRKLRTQETRCNEV
SHYKRLARSTLLLIPLGGIHYIVFAFGSEDAWEIQLFFELALGSFQGLVVAVLYCFLN
GEVQLEVQKKWQQWHLREFPLHPVASFSNSTKASHLEQSQGTCRTSII"
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16; Conser
                                                                                                                                                                                                                                                                       5-3, Yonbancho, Chiyoda-ku, Tokyo 102-0028 Japan For further information about this sequence, including its location and relationship to other sequences, please visit our sequence archive Web site (http://www-alis.tokyo.jst.go.jp/HGS/top. html) or send email to webmaster@www-alis.tokyo.jst.go.jp/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens genomic DNA, complete sequence.
APO00032
                                                                                                                                                                                                                                                                                                                                                                                                    This sequence is conducted by Kitasato University JST sequencing Leaboratory as a JST sequencing team. Principal Investigator:Yoshiyuki Sakaki Ph.D. Phone: +81-3-5449-5622, Fax: +81-3-5449-5445, phone: +81-3-5449-5622, Fax: +81-3-5449-5445, pakaki@hgc.ims.u-cokyo.ac.jp Sub-leader: Tadayoshi Shiba Ph.D., Masahira Hattori Ph.D. The sequence is submitted by:Human Genome Sequencing in ALIS project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Submitted. (11-MAY-1998) Masahira Hattori, Kitasato University, Department of Science, JST Sequencing Laboratory; Kitasato 1-15-1, Sagamihara 228, Japan (E-mail:hattori@hgc.ims.u-tokyo.ac.jp,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 (bases 1 to 100000)
Hattori,M., Ishii,K., Toyoda,A., Shiba,T. and Sakaki,Y.
Homo sapiens genomic DNA, chromosome 21q
Published Only in DataBase (1998)
2 (bases 1 to 100000)
Hattori,M., Ishii,K., Toyoda,A., Shiba,T. and Sakaki,Y.
Direct Submission
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                     Japan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tel:0427-78-9732, Fax:0427-78-9561)
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Yonbancho, Chiyoda-ku, Tokyo 102-0028 Jap
                                                                                                                                                              /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
/chromosome="21"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This sequence is conducted by Kitasato University JE Laboratory as a JST sequencing team. Principal Investigator: Yoshiyuki Sakaki Ph.D. Phone: +81-3-5449-5622, Fax: +81-3-5449-5445, sakaki@hgc.ims.u-tokyo.ac.jp Sub-leader: Tadayoshi Shiba Ph.D., Masahira Hattori sequence is submitted by Human Genome Sequencing in JST.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5-3, Yonbancho, Chiyoda-ku, Tokyo 102-0081 Japan For further information about this sequence, including is and relationship to other sequences, please visit our searchive Web site (http://www-alis.tokyo.jst.go.jp/HGS/) email to webmaster@www-alis.tokyo.jst.go.jp.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Submitted (15-APR-1999) Mika Hirakawa, Japan Science and Technology
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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5-3, Yonbancho, Chiyoda-ku, Tokyo 102-0081 Japan
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                                                                                                       /standard_name="D21S224"
/note="KM838/KM839;The location
site of PCR_primers."
                                  /note="SGC35207;The location of PCR primers."
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44275. .44383
                                                                                                                                                                                                                                                                                                                                                                                                                                          /mol_type="genomic_uni/db_xref="taxon:9606"
/chromosome="21"
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                                                                                           db_xref="GDB:182624"
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|1676. .11835
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/note="U77582;The location
)f PCR primers."
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                                                                                                                                                                                                                                                                        Submitted (10-MAY-1999) Masahira Hattori, The Institute of Physical and Chemical Research (RIKEN), Genomic Sciences Center (GSC); Kitasato Univ., 1-15-1 Kitasato, Sagamihara, Kanagawa 228-8555, Japan (E-mail:hattori@gsc.riken.go.jp, URL:http://hgp.gsc.riken.go.jp/, Tel:81-42-778-9923, Fax:81-42-778-9924)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ហ
                                                                                                                                  original/previous sequences. We determined the boundary between the insertion and genomic sequences experimentally, removed the insertion sequences, reconstituted the present data. The sequencing project is supported by Japan Science Technology Corporation (JST) and The Institute of Physical and Chemical Research (RIKEN).
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Hattori,M., Ishii,K., Toyoda,A., Taylor,T.D., Hong-Seog,P. Fujiyama,A., Yada,T., Totoki,Y. and Sakaki,Y.

Homo sapiens 2,083,744bp genomic DNA of 21q22.1 (REGION: D21s226-AML CLONE RANGE: Q76510-f3259)

Published Only in DataBase (1999)

2 (bases 1 to 10000)
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                                                                                                                                                                                                                                      E. coli transposon insertion: The present data does not contain E. coli transposon sequences which integrated in the
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site of PCR primers."
/db xref="CDB:6464271"
22604 c 22308 g 27882 t
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site of PCR primers."
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                                                  chromosome="21"
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Fax:81-42-778-9924)
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Submitted (08-NOV-1999, PROBLEM Sciences Center Coc., and Chemical Research (RIKEN), Genomic Sciences Center Coc., kitasato Univ., 1-15-1 Kitasato, Sagamihara, Kanagawa 228-8555, Japan (E-mail:hattori@gsc.riken.go.jp, Tel:81-42-778-9923,
                                                                                                                                                                                                                                                                  NOTE: This is a 'working draft' sequence. It currently consists of 13 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence
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Published Only in DataBase (1999)
2 (bases 1 to 109149)
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Center clone name: CMB9-2113

Center clone name: CMB9-2113

Sequencing vector: PCR products; 100% of reads

Sequencing vector: PCR products; 100% of reads

Chemistry: Dye-terminator ET-amersham; 100% of reads

Assembly program: Phrap; version 0.990329

Consensus quality: 99833 bases at least Q40

Consensus quality: 104437 bases at least Q30

Consensus quality: 104437 bases at least Q30

Consensus quality: 106696 bases at least Q20

Insert size: 107949; sum-of-contigs

Quality coverage: 4.90x in Q20 bases; sum-of-contigs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 109149)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Web site: http://hgp.gsc.riken.go.jp/
Contact: hattori@gsc.riken.go.jp
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b; Pred. No. 81;
0; Mismatches
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ri, The Institute of Physical Sciences Center (GSC);
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Sequence updated (26-May-2000).

NOTE: This is a 'working draft' sequence. It currently nonsists of 13 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence as soon as it is available and the accession number will be preserved.
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86790. .94162
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                                16;
         2 CTTAGGAGGAACAAGT 17
                                       Similarity
                                Conservative
                            80.0%; Pro
100.0%; Pro
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                                         Score 16;
Pred. No.
                               Mismatches
                                         DB
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                                                Length 109149;
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Search completed: August 15, 2003, 09:33:56
Job time : 555.75 secs

Run

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BW21593 AV821593
BJ273261 BJ273261
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BU25365 AGENCOURT
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             On Oct 3, 2000 this sequence version replaced gi:10575436. Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP/Gazdar
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Preparation: Life I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: The I.M.A.G.E. Consortium (Clone distribution: MGC clone distribution information can
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601658274R1 NIH_MGC_69
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                    Unpublished
                                                                                                                                                                                                                                                                                                     Homo sapiens (human)
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BUG9428 LL21n11197
BZ886748 CH240_269
AQ462877 HS 5212 A
BE40236 CSB00BD12
BQ608042 BRY 3943
AA422164 ZV91C01. r
BE288544 601094160
AG150407 Pan trog1
A1927904 wp03c07. x
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BG441695 GA Ea001
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BG442675 GA Ea001
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BH059230 RPCI-24-3
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CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by:Incyte Genomics, Inc.
Clone distribution: NCI-CGAP clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
plate: LLAM9279 row: p column: 08
High quality sequence stop: 672.
Location/Qualifiers
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1 (bases 1 to 817)
NIH-MGC http://mgc.nci.nih.gov/
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Plate: LLCM648 row: c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: cgapbs-r@mail.nih.gov
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Mammalia; Eutheria; Rodentia;
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/tissue_type="large cell carcinoma, undifferentiated"
/tissue_type="large cell carcinoma, undifferentiated"
/lab host="pH10B (phage resistant)"
/clone_lib="NIH MGC 69"
/cote="Organ: lung; Vector: pCMV-SPORT6; Site_1: NotI;
/note="Organ: lung; Vector: pCMV-SPORT6; Site_1: NotI;
/site_2: SalI; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.1 kb. Library constructed by Life
Technologies."
138 c 193 g 206 t
/lab_host="DH10B"
/clome_lib="MCI_CGAP_Lu30"
/clome_lib="MCI_CGAP_Lu30"
/note="Forgan: lung; Vector: pCMV-SPORT6; Site_1: NotI;
/note="Torgan: lung; Vector: pCMV-SPORT6; Site_1: NotI;
/notition="Tor
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Location/Qualifiers
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/mol_type="mRNA"
/strain="CZECH II"
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/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                        clone="IMAGE:4022647"
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Kim, N.S., Hahn, Y.,
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Location/Qualifiers
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Plate: 24 row: D column: 05
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Fax: +82-42-860-4409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Korea Research Institute of Bioscience & Biotechnology 52 Eoeun-dong Yuseong-gu, Daejeon 305-333, South Korea
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished
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                                                                                                                                                                         obtained cDNA vectors were used for transformation of competent cells E. coli Top10F' by electroporation method. The cDNA libraries constructed by this method are full-length enriched cDNA library."

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/cell_type="Epithelial"
/cell_line="SNU-601"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            lone="S9SNU601-24-D05"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     85.0%; br
100.0%; br
                                                                  80.0%; UT
100.0%; Pr/
0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 L_type="mRNA"
_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     lib="S9SNU601"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 17; pred. No.
                                                                                            Pred. No.
                                                                             Mismatches
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24;
                                                                                              DB 12;
65;
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                                                                                                                   Length 200;
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SOURCE
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ACCESSION
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BE755758
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                VERSION
KEYWORDS
                                                                                                                                                RESULT 5
Z42603
                                                                                                                                                                                                                                                                                                                                                                         ORIGIN
                                                                                                                                                                                                                                                                                                                                                                                            BASE COUNT
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                     SOURCE
                                                                         ACCESSION
                                                                                                             DEFINITION
                                    KEYWORDS
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MEDLINE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AUTHORS
                                                                                                                                                                                                                                                                                                                   Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                source
                                                                                                                                                                                                                           102
                                                                                                                                                                                                                                                                                                16;
Homo sapiens (human)
Homo sapiens
                                                                                                                                                                                                                                                           2 CTTAGGAGGAACAAGT 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Smith TPL
USDA, ARS, US Meat Animal Research Center
PO Box 166, Clay Center, NE 68933-0166, US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Smith, T.P.L., Grosse, W.M., Freking, B.A., Roberts, A.J., Stone, R.T. Casas, E., Wray, J.E., White, J., Cho, J., Fahrenkrug, S.C., Bennett (G.L., Heaton, M.F., Laegreid, W.W., Rohrer, G.A., Chitko-McKown, C. Chertea, G., Holt, I., Karamycheva, S., Liang, F., Quackenbush, J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bos taurus
Bos taurus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BE755758 203 bp mRNA linear
209682 MARC 2BOV Bos taurus cDNA 5', mRNA sequence.
BE755758
                                      EST
                                                                                                             Z42603 242 bp
HSC0PA091 normalized infant brain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BACKWARD: GTTTTCCCAGTCACGACG
Plate: 59 row: L column: 7
Seq primer: ATTTAGGTGACACTATAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Single pass sequencing. Bases called and alt trimmed with phred v0.980904.e. Vector identified by cross_match with the -minscore 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tel: 402 762 4366
Fax: 402 762 4390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence evaluation of four pooled-tissue normalized bovine cDNA libraries and construction of a gene index for cattle Genome Res. 11 (4), 626-630 (2001)
                                                     Z42603.1 GI:567353
                                                                                        c-Opa09, mRNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bovidae; Bovinae; Bos.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BE755758.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: smith@email.marc.usda.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Genome Res. 11 (4),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FORWARD: AGGAAACAGCTATGACCAT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -minmatch 12 options.
                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                            70
                                                                                                                                                                                                                                                                                                                                                                                            a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Bos taurus"
/mol_type="mRNA"
/db_xref="taxon:9913"
/tissue_type="pooled"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                  /note="Vector: PCWV SPORT6; Site 1: Not1; Site 2: Sal1; Library made from pooled tissue from testis, thymus, semitendonosus muscle, longissimus muscle, pancreas, adrenal, and endometrium."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ocation/Qualifiers
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                                                                                                                                                                                                                                                                                                                 80.0%;
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                                                                                                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                                                                                                 Score 16;
Pred. No.
                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                 65;
                                                                                                             cDNA Homo sapiens cDNA clone
                                                                                                                                mRNA
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AUTHORS
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KEYWORDS
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AU231688
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                                                                                                                                        TITLE
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1 (bases 1 to 242)

1 (bases 1 to 242)

1 (bases 1 to 242)

2 (bases 1 to 242)

3 (bases 1 to 242)

4 (bases 1 to 242)

4 (bases 1 to 242)

4 (bases 1, ..., to a silva, C..., D.

5 (bases 1, ..., Lore)

6 (bases 1, ..., Lore)

8 (bases 1, ..., Lore)

8 (bases 1, ..., Lore)

9 (bases 1, ..., Lore)

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AU231688 Cloned bovine mRNA sequence.
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Seq primer: (-21)M13_universal.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Genexpress-Genethon
Genethon Centre de recherche sur le Genome Humain
1,rue de l'Internationale, BP60 91002 EVRY Cedex, FRANCE
Graduate School of Agriculture
                                                                     Unpublished
                                                                                                 Analysis of expressed sequence tags from a cDNA library of somatic nuclear transfer-derived cloned bovine fetus
                                                                                                                                                                           Sasaki,Y.
                                                                                                                                                                                                1 (bases 1 to 350)
Taniguchi,Y., Lejukole,H.Y., Yamada,T.,
                                                                                                                                                                                                                                                                             Bovidae; Bovinae; Bos.
                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eute
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora;
                                                                                                                                                                                                                                                                                                                                                                           Bos taurus
                                                                                                                                                                                                                                                                                                                                                                                                                                              AU231688.1 GI:15719980
EST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Single read
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95277534
                                                                                                                                                                                                                                                                                                                                                                                                           Bos taurus (cow)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   7757816
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /tissue_type="total brain"
/dev_stage="3 months old"
/clone lib="normalized infant brain cDNA"
/clone lib="normalized infant brain cDNA"
/clone e="Organ: brain; vector: lafmid BA; Site_1: HindIII;
/note="Organ: brain; vector: lafmid BA; Site_1: HindIII;
/site_2: NotI; sex=Female; dev_stage=3 months old;
isolate=muscular atrophy patient; tissue type=total brain;
total mRNA was oligo-(dT) primed and directionally
cloned 5′ -> 3′ into the HindIII -> NotI sites of the
lafmid BA vector. Clone library from B. Soares, Psychiatry
Dept. Columbia University, USA. Normalization_method:
Bento Soares, P.N.A.S in press"
88 a 42 c 43 g 69 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ωı
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/db_xref="taxon:9606"
/clone="c-0pa09"
/sex="Female"
                                      Takahisa Yamada
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     organism="Homo sapiens"
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100.0%; Pred. No.
tive 0; Mismatc
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cDNA
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68;
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                                                                                                                                                                                                          Akagi,S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 242;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: est@watson.wustl.edu
Insert Size: 2215
High quality sequence stops: 284 Source: IMAGE Consortium, LLNL
High clone is available royalty-free through LLNL; contact the
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
IMAGE Consortium (info@image.llnl.gov)
INSERT Length: 2215
INSERT 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 358)

Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M., Holman
Hillier,L., Clark,N., Ed.,M., Lennon,G., Marra,M., Parsons,J.,
M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,J.,
Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Trevaskis,E., Waterston
,R., Williamson,A., Wohldmann,P. and Wilson,R.

The WashU-Merck EST Project
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This clone was obtained from a 3' end cDNA library.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sakyoku, Kitashirakawa, Kyoto, Kyoto 606-8502, Japan
Tel: 81-75-753-6323
Fax: 81-75-753-6340
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Fax: 314 286 1810
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/dev_stage="73 days post natal"
/lab_host="DH10B (ampicillin resistant)"
/clone_lib="Soares infant brain INIB"
/clone_lib="Soares infant brain INIB"
/note="Organ: whole brain; Vector: Lafmid BA; Site
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/clone_lib="Cloned bovine
70 c 83 g 92
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/db_xref="taxon:9913"
/clone="Cln595"
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/db_xref="GDB:406151"
/db_xref="taxon:9606"
/clone="IMAGE:33804"
                                                                                                                                                                                                                                                                                                                                                              'sex="female"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              organism="Homo sapiens"
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lty-free through LLNL ; contact the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Other_GSSs: RPCI-11-241P16.TV
Contact: Shaying Zhao, William Nierman,
Department of Eukaryotic Genomics
The Institute for Genomic Research
9712 Medical Center Dr., Rockville, MD
                            S
                                                                                                                                                                                                                                                                                                                                                                          library availability, please contact Pieter de Jong (pieter@dejong.med.buffalo.edu). Clones may be purchased from BACPAC Resources (http://bacpac.med.buffalo.edu/ordering) or from Research Genet cs (info@resgen.com). BAC end search page: http://www.tigr.org/tdb/humgen/bac_end_search/bac_end_search.html
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: hbe@tigr.org
Clones are derived from the human BAC library RPCI-11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Map Building
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Use of BAC End Sequences from Man Building
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Zhao, S., Adams, M.D., Nierman, W., Malek, J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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38: BAC ends
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                                                                                                                                                                                                                                  /organism="Homo sapiens"
/mol type="genomic DNA"
/db_xref="GDB:7592535"
/db_xref="taxon:9606"
/clone="RPCI-11-241P16"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         double-stranded cDNA was ligated to Hind III adaptors (Pharmacia), digested with Not I and directionally cloned into the Not I and Hind III sites of the Lafation BA vector. Library went through one round of normalization. Library constructed by Bento Soares and M.Fatima Bonaldo."

a 64 c 68 g 112 t 2 others
                                                                                                                                                      /clone_lib="RPCI-11"
/note="Vector: pBACe3.6; Site_1:
RPCI11 Human Male BAC Library"
                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                      cell_type="Lymphocytes"
                                                                                                                                                                                                                        sex="Male"
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Insert Length: 551 Std Error: 0.00
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Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Michael J. Br
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anat
Tumor Gene Index
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WC56dll.xl NCI_CGAP_Pr28 Homo sapiens cDNA clone IMAGE:2322645 similar to contains OFR.tl OFR OFR repetitive element ;, mRNA
                                                                                                       yc89c08.rl Soares infant brain IMAGE:23341 5', mRNA sequence.
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Seq primer: -40UP from Gibco
High quality sequence stop: 394.
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llarity 100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone lib="NCI_CGAP_pr28"
/note="Organ: prostate; Vector: pT773D-Pac (Pharmacia)
/note="Organ: prostate; Vector: pT773D-Pac (Pharmacia)
with a modified polylinker; Plasmid DNA from the
mormalized library NCI_CGAP_pr22 was prepared, and so
circles were made in vitro. Following HAP purification,
this DNA was used as tracer in a subtractive hybridization
this DNA was used as tracer in a full player than the property of the prope
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/mol_type="mRNA"
/db_xref="taxon:9606"
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lab_host="DH10B"
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Mismatches
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SOURCE

Homo sapiens (human)

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1 (bases 1 to 400)

Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M., Holman,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,J., Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P. and Wilson,R.

The WashU-Merck EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                   N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Insert Size: 2260
High quality sequence stops: 318 Source: IMAGE Consortium, LLNL This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Insert Length: 2260 Std Error: 0.00
Seq primer: M13RP1
                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 413)
                                                                                                                                                                                                                                 genomic survey sequence
AQ379270
                                                                                                                                                                                                                                                                     RPCI11-151H4.TJ RPCI-11 Homo
                Use of BAC End
Map Building
                                                                         Zhao, S., Adams M.D.,
                                                                                                                                                       Homo sapiens
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Location/Qualifiers
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Fax: 314 286 1810
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Washington University School of Medicine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished
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                                                                                                                                                                     Homo sapiens (human)
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                                                                                                                                                                                                                                                                                                                                                                                                                 CTTAGGAGGAACAAGT 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    double-stranded cDNA was ligated to Hind III adaptors (Pharmacia), digested with Not I and directionally cloned into the Not I and Hind III sites of the Lafand BA vector. Library went through one round of normalization. Library constructed by Bento Soares and M.Fatima Bonaldo."

77 c 72 g 127 t 1 others
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Homo sapiens"
/mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /dev_stage="73 days post natal"
/lab_host="DH10B (ampicillin resistant)"
/clone_llb="Soares infant brain INIB"
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/clone="IMAGE:23341"
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                                  Sequences from Library RPCI-11 for Sequence-Ready
                                                                       Nierman, W., Malek, J.,
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                                                                                                                                                                                                                                                                        genomic clone
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                                                                         de Jong, P. and Venter
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AW769407
                                             FEATURES
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                                                                                                                                                        Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can found through the I.M.A.G.E. Consortium/LLNL, send email to:
                                                                                                                                                                                                                                    Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Chris Moskaluk, M.D., Ph.D., Michael R.
Tissue Procurement: Chris Moskaluk, M.D., Ph.D. Life
Emmert-Buck, M.D., Ph.D. CDNA Library Preparation: Life
Technologies, Inc. CDNA Library Arrayed by: Christa Prange, The
I.M.A.G.E. Consortium DNA Sequencing by: Washington University
                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (baes 1 to 431) NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap. NCI-CGAP institute, Cancer Genome Anatomy Project (CGAP), mational Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AW769407

431 bp mRNA linear EST 04-MAY-2000 hl63g12.x1 NCI_CGAP_Kid13 Homo sapiens cDNA clone IMAGE:3005926 3' similar to contains OFR.bl OFR repetitive element ;, mRNA sequence.
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Clones are derived from the human BAC library RPCI-11. For BAC

Clones are derived from the human BAC library RPCI-11. For BAC

Clones are derived from the human BAC library availability, please contact Pieter de Jong

(pieter@dejong.med.buffalo.edu). Clones may be purchased from BACPAC Resources (http://bacpac.med.buffalo.edu/ordering) or from Research Genetics (info@resegen.com). BAC end search page:

http://www.tigr.org/tdb/humgen/bac_end_search/bac_end_search.html
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Contact: Shaying Zhao, William Nierman,
Department of Eukaryotic Genomics
The Institute for Genomic Research
The Institute for Genomic Research
                                                               Possible reversed clone: polyT not
Seq primer: -40UP from Gibco
High quality sequence stop: 396.
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Fax: 301 838 0208
                                                                                                                                                                                                                                                                                                                                                                 Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                   Tumor Gene Index
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
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                                                                                                                                           info@image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens (human)
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RPCII1 Human Male BAC Library"
92 c 92 g 104 t 1 others
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/db_xref="taxon:9606"
/clone="RPCI-11-151H4"
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/mol_type="genomic DNA"
/organism="Homo sapiens
                                           Location/Qualifiers
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76;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Grossman,A., Davies,J., Federspiel,N., Harris,E., Lefebvre,P., McDermott,J.P., Silflow,C., Stern,D. and Surzycki,R. Analyses of the Chlamydomonas reinhardtii Genome: A Model, Unicellular System for Analyzing Gene Function and Regulation in Vascular Plants; project phase 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BE227333 443 bp mRNA lin 894030E10.x3 C. reinhardtii CC-1690, normalize Chlamydomonas reinhardtii cDNA, mRNA sequence.
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Chlamydomonas reinhardtii
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Durham, NC 277408-1000, USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Vascular Plants; project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Viridiplantae; Chlorophyta; Chlorophyceae; Volvocales; Chlamydomonadaceae; Chlamydomonas.
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919 613 8177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chlamy@duke.edu.
                                                                                                                                                                                 /note="Vector: pBluescript II SK-; Site_1: EcoRI; Site_2: XhoI; This library, constructed by John Davies and Jeffrey McDermott, combines cDNAs from CCC-1690 cells grown to mid-log phase in TAP (acetate-containing) medium in the light, TAP medium in the dark, HS (minimal) medium in ambient levels of CO2 and HS medium bubbled with 5% CO2. POlyA mRNA was purified from each sample, pooled and cDNA synthesized. The cDNA was directionally cloned into lambda SAP II (Stratagene) in the EcoRI (5') and XhoI (3') sites.
                                                           ZAP II (Stratagene) in the EcoRI (5') and XhoI (3') sites. pBluescript II SK- plasmids were excised from the lambda ZAP clones by superinfection with ExAssist (Stratagene) phage. The library was normalized using method 4 described in Bonaldo et al (1996) Genome Research 6: 791-806."

129 c 139 g 86 t
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/note="Organ: kidney; Vector: pCMV-SPORT6; Site_1: SalI;
Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Library constructed by Life Technologies.
98 c 103 g 99 t
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/lab_host="DH10B"
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/tlasue_type="2 pooled Wilms' tumors, one primary and one
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
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/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                  /mol_type="mRNA"
/strain="CC-1690 wild type mt+ 21gr"
/db_xref="taxon:3055"
                                                                                                                                                                                                                                                                                                                                                                                                                         clone_lib="C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          organism="Chlamydomonas reinhardtii
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  80.0%;
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Score 16;
                                                                                                                                                                                                                                                                                                                                                                                                                            reinhardtii CC-1690, normalized, Lambda Zap
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77;
  DB 10;
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abda Zap II
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COMMENT

ORIGIN

prepare full-length cD genes. Genome Res. 10

Genome Sequences Mamm.

Normalization and subtraction of cap-trapper-selected cDNAs to prepare full-length cDNA libraries for rapid discovery of new yenes. Genome Res. 10 (10), 1617-1630 (2000)

integrated sequence analysis (RISA)

밁 8 ORIGIN

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REFERENCE
AUTHORS
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MEDLINE
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                                                                                             Email: genome-res@gsc.riken.go.jp,
URL:http://genome.gsc.riken.go.jp/
Alzawa,K., Akimura,T., Arakawa,T., Carninci,P., Fukuda,S., Hirozane
Alzawa,K., Akimura,T., Arakawa,T., Carninci,P., Fukuda,S., Hirozane
,T., Imotani,K., Ishii,Y., Itoh,M., Kawai,J., Konno,H., Miyazaki,A.,
,T., Murata,M., Nakamura,M., Nomura,K., Numazaki,R., Ohno,M., Sakai,K.,
,Sakazume,N., Sasaki,D., Sato,K., Shibata,K., Shiraki,T., Tagami,
,Sakazume,N., Sasaki,D., Sato,K., Shibata,K., Shiraki,T., Tagami,
,M., Waki,K., Watahiki,A., Muramatsu,M. and Hayashizaki,Y. Direct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Laboratory for Genome Exploration Research Group, RIKEN Genomic Sciences Center(GSC), Yokohama Institute
The Institute of Physical and Chemical Research (RIKEN)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nature 420,
22354683
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Analysis of the mouse transcriptome based on functional annotation of 60,770 full-length cDNAs
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                                                                                                                                                                                                                                                                                                                                                                                                           1-7-22 Suehiro-cho,
Tel: 81-45-503-9222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12466851
                                                                                                                                                                                                                                                                                                                                                                               Fax: 81-45-503-9216
Computational Analysis of Full-Length Mouse cDNAs Compared with 
luman Genome Sequences Mamm. Genome. 12, 673-677 (2001)
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cDNA clone 1920006J23 3', mRNA sequence.
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JOURNAL
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Best Local
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16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Computer-based methods for the mouse full-length cDNA encyclopedia: real-time sequence clustering for construction of a nonredundant cDNA library. Genome Res. 11 (2), 281-289 (2001) cDNA library was prepared and sequenced in Mouse Genome Encyclopedia Project of Genome Exploration Research Group in Riken Genomic Sciences Center and Genome Science Laboratory in RIKEN. Division of Experimental Animal Research in Riken contributed to
                                                                                                                                                                                                                                                                                     Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. I Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                   Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M., Holman,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,J. Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P. and Wilson,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    yg32f03.r1 Soares infant brain 1NIB Homo sapiens cDNA clone IMAGE:33987 5', mRNA sequence.
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Please visit our web site (http://genome.gsc.riken.go.jp) for
                                                                                                                                                                         High quality sequence stops: 349 Source: IMAGE Consortium, LLNL This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               R24326.1 GI:739214
EST.
                                                                                                                                                                                                                                                                Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                 Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        further details.
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                                                                                                                                                       primer: M13RP1
                                                                                                                                  quality sequence stop: 349.
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/dev stage="16 days embryo"
/clone_lib="RIKEN full-length enriched,
kidney"
/organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="GDB:406334"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /mol_type="mRNA"
/strain="C57BL/6J"
                                                                                                              ocation/Qualifiers
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'clone="1920006J23"
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                                                                                                                                                                                                                                                                      Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=MR3&t2=MR3-HN0052-
271200-001-b01&t3=2000-12-27&t4=1)
Seq primer: puc 18 forward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          l (bases 1 to 487)
Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R.,
Dias Neto,E., Garcia Correa,R., Bordin,S., Costa,F.F.,
Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F.,
Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H.,
Brunstein,A., deOliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare
M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rua Prof. Antonio Prudente 109,
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Tel: +55-11-2704922
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                                                                                                                                                                                                      quality sequence start: 32 quality sequence stop: 463. Location/Qualifiers
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87 c 86 g 148 t 1 others
/clone_lib="HN0052"
/clone_Togan: head_normal; Vector: puc18; Site_1: SmaI;
/note="Organ: head_normal; Vector: puc18; Site_1: SmaI;
Site_2: SmaI; A mini-library was made by cloning products
derived from ORESTES PCR (U.S. Letters Patent application
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/db_xref="taxon:9606"
/dev_stage="Adult"
                                                                                                                                                           organism="Homo sapiens"
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1 (bases 1 to 494)

Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M., Holman, M., Hultman, M., Kucaba, T., Le, M., Lennon, G., Marra, M., Parsons, J., Rifkin, L., Rohlfing, T., Soares, M., Tan, F., Trevaskis, E., Waterston, R., Williamson, A., Wohldmann, P. and Wilson, R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: est@watson.wustl.edu
Insert Size: 2185
High quality sequence stops: 325 Source: IMAGE Consortium, LLNL
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Insert Length: 2185 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Washington University School of Medicine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               yg32h03.rl Soares infant brain INIB Homo sapiens cDNA clone IMAGE.34200 5', mRNA sequence.
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profiles into the pUC 18 vector. Reverse transcription
tissue mRNA and cDNA amplification were performed under
low stringency conditions."
102 c 108 g 112 t
                                                                                                                         I; Site 2: Hind III; 1st strand cDNA was primed with a Not I - oligo(dT) primer [5]
AACTGGAAGAATTGTGGAGGAATTGTTTTTTTTTTT 3'];
double-stranded cDNA was ligated to Hind III adaptors (Pharmacia), digested with Not I and directionally cloned into the Not I and Hind III sites of the Lafnid BA vector. Library went through one round of normalization. Library
                                                                                   constructed by Bento Soares and M.Fatima Bonaldo.
1 90 c 90 g 159 t 4 others
                                                                                                                                                                                                                                                                              /dev stage="73 days post natal"
/lab_host="DH10B (ampicillin resistant)"
/clone lib="Soares infant brain INIB"
/note="Organ: whole brain; Vector: Lafmid BA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Homo sapiens"
/mol_type="mRNA"
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                                                                                                                                                                                                                                                                                                                                                                      sex="female"
                                                                                                                                                                                                                                                                                                                                                                                          clone="IMAGE:34200"
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db_xref="taxon:9606"
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                                                                                                                                                                                                                                                        or: Lafmid BA; Site_1: Not cDNA was primed with a Not
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: genome-res@gsc.riken.go.jp,

URL.http://genome.gsc.riken.go.jp/

Carninci,F., Shibata,Y., Hayatsu,N., Sugahara,Y., Shibata,K., Itoh
,M., Konno,H., Okazaki,Y., Muramatsu,M. and Hayashizaki,Y.

Normalization and subtraction of cap-trapper-selected cDNAs to
prepare full-length cDNA libraries for rapid discovery of new
genes. Genome Res. 10 (10), 157-1630 (2000)

wagi,K., Fujiwake,S., Inoue,K., Togawa,Y., Izawa,M., Ohara,E.,
Watahiki,M., Yoneda,Y., Ishikawa,T., Ozawa,K., Tanaka,T., Matsuura
,S., Kawai,J., Okazaki,Y., Muramatsu,M., Inoue,Y., Kira,A. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Yoshihide Hayashizaki
Laboratory for Genome Exploration Research Group, RIKEN
Sciences Center(SC), Yokohama Institute
The Institute of Physical and Chemical Research (RIKEN)
1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-
Tel: 81-45-503-9222
Fax: 81-45-503-9216
                                                                                                                                                                                                                                                                                                                                                                                                                    encyclopedia: real-time sequence clustering for construction of nonredundant cDNA library. Genome Res. . 11 (2), 281-289 (2001) Please visit our web site (http://genome.gsc.riken.go.jp) for further details.
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BB693705
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Computer-based methods for the mouse full-length cDNA
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/clone lib="RIKEN full-length enriched, sympathetic ganglion"
                                                                                                                                                                                                                                                                    /organism="Mus musculus"
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                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                              'db_xref="taxon:10090"
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                                                                                                                                                         sex="mixed"
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                                                            Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml1.pl?ti=PMO&t2=PMO-HT1166-
150201-002-c08&t3=2001-02-15&t4=1)
Seq primer: puc 18 forward
High quality sequence start: 10
High quality sequence stop: 516.
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Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R.,
Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F.,
Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H.,
Brunstein,A., deOliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare
,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
                                                                                                                                                                                                                                                                                                                                             Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
                                                                                                                                                                                                                                                                                                                              Rua Prof. Antonio Prudente 109, 4 andar,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Shotgun sequencing of the human transcriptome with ORF expressed
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PM0-HT1166-150201-002-c08
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sequence tags
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             assistance we gratefully acknowledge."
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organism="Homo sapiens"
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3 HT1166 Homo sapiens
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SOURCE KEYWORDS VERSION

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1 (bases 1 to 524)
Yang, R.-Z., Shuldiner, A.
EST analysis of human ad
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1001704 Human Fat Cell 5'-Stretch Plus cDNA Library Homo sapiens cDNA 5', mRNA sequence.
CA405555
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                                                                                                                                                                                                                                                                                                                        BACKWARD: AATACGACTCACTATAGGGCGAATTGG
Seq primer: GTTGGTACCCGGGAATTC.
                                                                                                                                                                                                                                                                                                                                                                                                              Tel: 410 706 1672
Fax: 410 706 1622
                                                                                                                                                                                                                                                                                                                                                                                                                                                 University of Maryland 660 Redwood St, HH497,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Division of Endocrinology, Diabetes and Nutrition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Gong Da-Wei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                            Email: dgong@medicine.umaryland.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens (human)
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                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note="Organ: head neck; Vector: puc18; Site_1: SmaI; Site_2: SmaI, A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under low stringency conditions."
                                                                                                                                             /sex="Male and Female"
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/clone Ii="Human Fat Cell 5'-Stretch Plus
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/dev_stage="Adult"
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                                                                                                     zc94e08.rl Pancreatic
5', mRNA sequence.
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Dias Neto, E., Garcia Correa, R., Verjovski-Almeida, S., Briones, M.R., Nagai, M.A., da Silva, W. Jr., Zago, M.A., Bordin, S., Costa, F.F., Nagai, M.A., da Silva, W. Jr., Zago, M.A., Bafa, G.S., Simpson, D.H., Goldman, G.H., Carvalho, A.F., Matsukuma, A., Bafa, G.S., Simpson, D.H., Brunstein, A., deOliveira, P.S., Bucher, P., Jongeneel, C.V., O'Hare, M.J., Soares, F., Brentani, R.R., Reis, L.F., de Souza, S.J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (http://www.ludwig.org.br/scripts/gethtml2.pl7t1=CM0&t2=CM0-HT0179-051099-064-g10&t3=1999-10-05&t4=1)
Seq_primer: puc 18 forward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
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                                 Homo sapiens
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This sequence was derived from the FAPESP/LICR Human
Project. This entry can be seen in the following URL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Simpson, A.J.
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Eukaryota; Metazoa;
Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Shotgun sequencing of the human transcriptome with ORF expressed
                                                   Homo sapiens (human)
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                     /note="Organ: head_neck; Vector: puc18; Site_1: SmaI; Site_2: SmaI, A mini-library was made by cloning products derived from ORSTES PGR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under low stringency conditions."

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/mol_type="mRNA"
/db_xref="taxon:9606"
/dev_stage="Adult"
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81;
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Contact: Wilson RK
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Ta01_05d02_C
Ta01_AAFC_ECORC_Fusarium_graminearum_inoculated_wheat_heads
Triticum_aestivum_cDNA_clone_Ta01_05d02, mRNA_sequence.
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Eastern Cereal and Oilseed Research Centre
Agriculture and Agri-food Canada
Neatby Bldg., Central Experimental Farm, Ottawa,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Viridiplantae; Streptophyta; Em
Spermatophyta; Magnoliophyta; Liliopsida;
; Triticeae; Triticum.
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                                                                                                                                      Contact: Ouellet, Therese
                                                                                                                                                                                                                                                                                                                                                                                                           Ouellet, T., Dan, H., Koul, A.,
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                                                                                                                                                                                                                                                                                                                                                   Harris, L.J., Hattori, J.I.,
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/lab_host="SOLR cells (kanamycin resistant)"
/clone_lib="Pancreatic Islet"
/clone_tib="Pancreatic Islet"
/note="Organ: pancreas; Vector: pBluescript SK-; Site 1:
/note="Torgan: pancreas; Vector: pBluescript SK-; Site 1:
/conte="Torgan: pancreas; Vector: pBluescript SK-; Site 1:
/rote="Torgan: pancreas; Vector: pBluescript SK-; Sit
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         K1A 0C6,
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RESULT 24
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Best Local
                                                                                                          Clone from S. Wiemann, Molecular Genome Analysis, German Ca
Research Center (DKF2); Email s.wiemann@dkfz-heidelberg.de;
sequenced by AGOWA (Berlin/Germany) within the cDNA sequenc
consortium of the German Genome Project.
                                                                                                                                                                                                                                                                                                                                                                                                                  09-MAY-2003
09-MAY-2003
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source
                         Key
                                                                     Please
                                                                               No s1 sequence available.
This clone (DKFZp779F1664) is available at the RZPD in Berlin.
                                                                                                                                                               This is the 5' sequence of the clone insert
                                                                                                                                                                                            MIPS,
                                                                                                                                                                                                          Submitted
                                                                                                                                                                                                                                                 Wambutt
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                                                                                                                                                                                                                                                                                                                                   Homo sapiens (human)
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                                                   14059 Berlin-Charlottenburg,
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Ingolstaedter Landstr.1, D-85764 Neuherberg, GERMANY
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                                                  the RZPD: Ressourcenzentrum, Heubnerweg 6, harlottenburg, GERMANY; Email: clone@rzpd.de
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                                                                                                                                                                                                                                                                                                                                                             sequence tag
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a 157 c 132 g 140 t
                         Location/Qualifiers
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/clone_lib="Ta01_AAFC_ECORC_Fusarium_graminearum_inoculate
1_wheat_heads"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         organism="Triticum aestivum"
(mol type="mRNA"
(cultivar="Frontana"
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clone="Ta01_05d02"
tissue_type="heads"
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75,
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Last updated, Version
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                                                                                                                       cDNA sequencing
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RESULT 26
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16; Conserv
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AU137185
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16; Conserv
BI156614
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Tel: 81-438-52-3975
Fax: 81-438-52-3986
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                                                                                                                                                                                                                                                                                         Research Institute; cDNA library construction: Virology, Institute of Medical Science, Univers Helix Research Institute.
                                                                                                                                                                                                                                                                                                                                                                         Genomics Laboration:
Helix Research Institute
Kisarazu, Chiba
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ota,T., 'Nishikawa,T., Suzuki,Y., Yamamoto,J., Wakamatsu,A., Nakam
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AU137185.1
EST.
                                                                                                                                                                                                                                                                                                                           Email: genomics@hri.co.jp
HRI human cDNA project; 5'- & 3'-end one pass sequencing: Helix
                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Takao Isogai
                                                                                                                                                                                                                                                                                                                                                                                                                            Unpublished
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                                                                          CTTAGGAGGAACAAGT 17
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                                                                                                                                                                    200
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ilarity 100.0%;
Conservative
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/mol_type="mRNA"
/organism="Homo sapiens"
/clone="DKFZp779F1664"
/clone=lib="779 (synonym: hnccl). Vector pSport1_Sfi; host DH10B; stees Sfila + SfilB"
/dev_stage="fetal"
                                                                                                                                                                                                                  /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="PLACE1005960"
                                                                                                                                                                 /note="Vector: pME18SFL3"
124 c 92 g 196
                                                                                                                                                                                           /tissue_type="placenta"
/clone_lib="PLACE1"
                                                                                                                                                                                                                                                                              ocation/Qualifiers
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82;
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83
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ura,Y., Nagai,T., Sugano,S. and
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PLACE1005960
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                                                                                                                                   Arabidopsis thaliana (thale cress)
Arabidopsis thaliana
Arabidopsis thaliana
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Plate: LLAM11166 row: k column: 02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    602921206F1 NCI_CGAP_Mam3 Mus musculus cDNA clone IMAGE:5061625 5',
Large scale analysis of Arabidopsis full-length cDNA (2002b) Unpublished
                                                                         Seki,M., Narusaka,M., Ishida,J., Kamiya,A., Satou,M., Nakajima,M., Oono,Y., Sakurai,T., Carninci,P., Kawai,J., Itoh,M., Ishii,Y.,
                                                                                                                                                                                                                                                                              mRNA sequence.
AV821593
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NIH-MGC_http://mgc.nci.nih.gov/.
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                                           and Shinozaki, K.
                                                         Arakawa, T., Shibata, K., Shinagawa, A., Muramatsu, M., Hayashizaki, Y.
                                                                                                                                                                                                                                                       AV821593.1 GI:19863621
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /tissue_type="tumor, gross tissue"
/dev_stage="10 months"
/lab_host="DH10B"
/lab_host="HJH10B"
/clone_lib="NCI_CGAP_Mam3"
/clone_stage="NCI_CGAP_Mam3"
/note="Organ: mammary; Vector: pcWV-SPORT6; Site_1: Sall;
Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT.
Library constructed by Life Technologies. Investigators
providing samples: Lothar Hennighausen/Chu-Xia Deng, NIH
Reference for transgenic model: Xu et al., Nature Genetics
22, 37-43 (1999)."
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/strain="129,C57BL/6J,FVB/N"
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100.0%; Pred. No.
Live 0; Mismatc
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83;
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COMMENT
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An Arabidopsis full-length cDNA library was constructed essentially as reported previously (Seki et al., 1998).This clone is in a modified pBluescript vector as a SstI/XhoI insert. Please visit our web site (http://www.gsc.riken.go.jp/e/plant/index _e.html) for
                                                                                                                                                                                                                                                                                                                       Center For Genetic Resource Information
National Institute of Genetics
1111 Yata, Mishima, Shizuoka 411-8540, Japan
                                                                                                                                                                                                                                                                                                                                                                                                                  1 (bases 1 to 668)
Ogihara, Y. and Murai, K.
Expressed genes in Triticum aestivum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BJ273261 Y. Ogihara unpublished aestivum cDNA clone whohl6g03 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 . Similarity 16; Conser
                                                                                                                                                                                                                                                                                   Tel: 81-559-81-6856
Fax: 81-559-81-6855
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Spermatophyta; Magnoliophyta; ; Triticeae; Triticum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Triticum aestivum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tel: 81-298-36-4359
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RIKEN Genomic Sciences Center
3-1-1 Koyadai, Tsukuba, Ibaraki 305-0074, Japan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Motoaki Seki
                                                                                                                                                                                                                                                                                                                                                                                Contact: Tadasu Shin-i
                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Viridiplantae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Triticum aestivum (bread wheat)
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                                                                                                                                                                                                                                           tshini@genes.nig.ac.jp
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /clone_lib="RAFL4"
/note="Site_1: SstI; Site_2: XhoI; subjected
cold-treated(1,2,5,10,24 hr)"
132 c 140 g 201 t
                                                   /tissue_type="pistil at heading date"
/dev_stage="Feekes' scale 10.5"
/clone_lib="Y. Ogihara unpublished cDNA library, Wh_oh"
185 c 148 g 163 t
                                                                                                                                /mol type="mRNA"
/cultivar="Chinese Spring"
/db xref="taxon:4565"
/clone="whoh16g03"
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                                                                                                                                                                                                        organism="Triticum aestivum"
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a; Poales; Poaceae; Pooidea
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ACCESSION
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BU635192
LOCUS
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Arabidopsis thaliana Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosid; eurosids II; Brassicales; Brassicaceae; Arabidopsis.

1 (bases I to 740)
Lundsgaard,M., Emmersen,J., Nielsen,K.L., Wilson,I., Somerville,S. and Welinder,K.G.
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                                                                                                                                                                                                                                                                                                                                                        16;
                                                                                                                                                                                                    BU635192 740 bp mRNA linear ES 003D06 Infected Arabidopsis Leaf Arabidopsis thaliana
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Scala, S., Carels, N., Falciatore, A., Chiusano, M.L. and Bowler, C. Genome properties of the diatom Phaeodactylum tricornutum Plant Physiol. 129 (3), 993-1002 (2002)
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BQ539794.1 GI:21395364
EST.
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Phaeodactylum tricornutum
Eukaryota, stramenopiles, Bacillariophyta,
Bacillariophycidae, Naviculales, Phaeodacty
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PTAM0102 Phaeodactylum tricornutum
tricornutum cDNA 5', mRNA sequence.
                                                                                                                        Arabidopsis thaliana (thale cress)
                                                                                                                                                      BU635192.1 GI:23302447
                                                                                                                                                                                    sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: chris@alpha.szn.it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Bowler C
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llarity 100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Phaeodactylum
/mol_type="mRNA"
/db xref="taxon:2850"
/cell_line="CCMP632"
                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone_lib="Phaeodactylum tricornutum Uni-Zap XR"
/note="Vector: Uni-Zap XR vector; Site_1: Eco RI; Site_2:
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                                                                                                                                                                                                                 CDNA Library Preparation: Ling Hong/Rubin Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLCM1842 row: e column: 15
High quality sequence stop: 674.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 745)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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Aalborg Universitet
Sohngaardsholmsvej 49, 9000 Aalborg,
                                                                                                                                                                                                                                                                                                                                                                                  Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
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Fax: +45 98141808
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                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert Strausberg,
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/clone="IMAGE:5022326"
/tissue_type="rhabdomyosarcoma"
/lab_host="DH10B (phage-resistant)"
/clone_lib="NH1M_MGC_17"
/clone_tib="NH1M_MGC_17"
/note="Organ: muscle; Vector: pOTB7; Site_1:
Site_2: XhoI; cDNA made by oligo-dT priming.
Directionally cloned into EcoRI/XhoI sites us
following 5' adaptor: GGCACGAG(G). Size-sele
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library of Arabidopsis and E. cichoracearum infecte
from three weeks old Arabidopsis plants. Plants wer
                                                                                                                                               /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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/strain="Columbia"
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'clone_lib="Infected Arabidopsis Leaf"
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National Institutes of Health, W
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 http://image.llnl.gov
Plate: LLAM9567 row: f column: 19
High quality sequence stop: 676.
Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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mRNA sequence.
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Clone distribution: MGC clone distribution information can
cound through the I.M.A.G.E. Consortium/LLNL at:
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a 244 c 204 g 146 t
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                                                                                                                                                                                                                                                Technologies. "
. 169 c 161 g
                                                                                                                                                                                                                                                                     Average insert size 1.8 kb. 
Technologies. "
                                                                                                                                                                                                                                                                                                                          tissue_type="adenocarcinoma"

lab_host="DH10B (phage-resistant)"

clone_lib="NIH_MGC_65"
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IMAGE:4992843 5',
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                                                                                                                     Contact: Robert Strausberg, Email: cgapbs-r@mail.nih.go. Tissue Procurement: ATCC
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1 (bases 1 to 879)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)
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AGENCOURT 7940818 NIH MGC 71
5', mRNA sequence.
Bill 77000
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection
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BI088175.1 GI:14506505
                                                                                                                                                                          Unpublished
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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             found through the I.M.A.G.E. Consortium/LLNL http://image.llnl.gov
                                              cDNA Library Preparation: Life Technologies, Inc. CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Agencourt Bioscience Corporation Clone distribution: MGC clone distribution information can
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/note="Organ: cervix; Vector: pCMV-SPORT6; Si
Site 2: Sall; Cloned unidirectionally. Prime
Average insert size 1.5 kb. Library prepared
Technologies."
165 c 151 g 275 t
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/lab_host="DH108"
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cell_line="MGC36"
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/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                   found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov plate: LLMM10628 row: h column: 08 High quality sequence stop: 753.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tissue Procurement: James Cleaver, M.D.

CDNA Library Preparation: Life Technologies, Inc.

CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA

Sequencing by: Incyte Genomics, Inc.

Clone distribution: MGC clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 884)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)
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884 bp mRNA linear EST 15-MAY-2001
602631028F1 NCI_CGAP_Skn3 Homo sapiens cDNA clone IMAGE:4776247 5',
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Location/Qualifiers
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Average insert size 1.5kb. Library constructed by Life
Technologies. Note: this is a NCI_CGAP Library."
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159 c 143 g 297 t
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